

STIC Search Report Biotech-Chem Library

STIC Database Tracking Number: 140014

TO: Andrew D Kosar

Location: rem/3c04/3c18

Art Unit: 1654

Tuesday, December 14, 2004

Case Serial Number: 10/068905

From: Edward Hart

Location: Biotech-Chem Library

REM-1A55

Phone: 571-272-2512

edward.hart@uspto.gov

Search Notes

Examiner Kosar,

Here are the results of the search you requested.

Please feel free to contact me if you have any questions.

Edward Hart



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STIC-Biotech/ChemLib

140014

From:

Kosar, Andrew

Sent:

Friday, December 10, 2004 10:44 AM

To:

STIC-Biotech/ChemLib

Subject: 10/068,905

Please search in 10/068,905 SEQ ID NO: 5, 6, 7, 8, and 9. They are cyclic tetrapeptides. thank you, Andrew Kosar

Andrew D. Kosar, Ph.D. Art Unit 1654 REMSEN 3C04 Mail REMSEN 3C18

(571)272-0913

DEC 10 2005

TOTAL CHERT DIVINE

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Maximum :
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AAR71313
AAR5533
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17.936 Million cell updates/sec
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Compugen Ltd.
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Aap71313 Peptide c
Aar55333 Inhibitor
Aar75499 C5a recep
Abp55713 B15 class
Abp55796 B15 class
                                             Abp67652 Human (
Abp67653 Human (
Abp67653 Human (
Abp67653 Erbs re
Abj18638 Erbs re
Abj99243 Peptide
Adj81984 Thermus
Adl19858 Human 1
Adl9859 Human 1
Adl98516 Human 1
Adl98518 Human 1
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Adl98678 Human 1
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Aar43432
                                                                                                                                                                                                                                                              Description
       Human leu
Human leu
Antibacte
Antibacte
Ro/SSA ep
Second ge
Synthetic
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Human CD6
Human CD6
Caspase C
Caspase C
FibB rece
Peptide P
Formus b
Human leu
Human leu
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25-MAR-2003 03-OCT-2002 21-AUG-1991

(revised)
(revised)
(first entry)

AAP61656;

AAP61656 standard; protein; 4

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Sequence of peptide which inhibits kinase activity and mammalian cell

 $\begin{tabular}{ll} $cyclic-nucleotide independent \\ $growth. \end{tabular}$

Cell growth

inhibitor; tumour cell

growth inhibitor

45	44	43	42	41	40	39	38	37	36	35	34	w	32	31	30	29	28	27	t
13	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13	13	
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Abp59244	Abj18639	Abb75621	Aae16333	Abp67642	Abp67641	Aab66249	Aab66243	Aab08397	Aay68375	Aay68369	Aay44679	Aay43367	Aay43373	Aaw95236	Aaw95230	Aay41826	Aay41820	Aar72699	
Peptide		Anti-RANK	N-termina	Human CD6	H		B pertuss	Peptide d	Pertussi		Hydropero	Pertussis	Pertussis	PT toxin	ΡŢ	Pertussis		Kappa-cas	

ALIGNMENTS

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RESULT 1
AAP61656
AXX AAP6
AXX 25-M
AXX 6-M
AXX Call
AXX Call
AXX AP6
AXX 16-M
AXX 1
                         The cpds. of the invention inhibit protein phosphorylation. The inventors claim a process for inhibiting the growth of tumour cells in a medium which comprises contacting the cells with a cpd. of formula (AAPs1614-P61661) or a physiologically acceptable salt. (Updated on 03-OCT-2002 to add missing OS field.) (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                      Claim 1; Col 4; 9pp;
                                                                                                                                                                                                                                                                                                    Inhibition of tumour cell growth - halo:methyl ketone(s).
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RESULT 3
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AC AAR5
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                             25-MAR-2003
30-JAN-1995
                                                                                                                                                                                                 Sequence 4 AA;
                                                                                                                                                                                                                           This peptide is useful as part of a peptide/halo-methyl ketone cpd., for treating picornavirus, egpolio- or rhimovirus infections. It inhibits the processing of picornavirus capsid proteins by virus encoded proteases. See AAP71301-12. See also US4652552. (Updated on 25-MAR-2003 to correct
                                                                                                                                                                                                                                                                                          Treating picorna-virus infection with peptide halo:methyl ketone esp. for treating polio virus and rhino virus infections.
                                                                                                                                                                                                                                                                                                                         ildm
           Inhibitor of tyrosine kinase
                                                         AAR55333;
                                                                          AAR55333 standard;
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                                                                                                                                                                                                                                                                                                                                           Kettner CA,
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15-MAY-1991
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RESULT 4
AAR75499
ID AAR7
XX AAR7
AC AAR7
XY 25-W
DT 13-W
XX C5a,
XX C5a,
XX W09:
XX W09:
XX W09:
XX 16-1
XX 16-1
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XX (C1)
XX (C1)
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Matches 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 4 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New peptide inhibitors of protein tyrosine kinase - contain D-Tyr or tetrafluoro-Tyr residue, for treating proliferative diseases, viral
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27-OCT-1993;
                                             06-DEC-1993;
                                                                          16-NOV-1994;
                                                                                                       15-JUN-1995.
                                                                                                                                     WO9516033-A1
                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                 C5a;
                                                                                                                                                                                                                            C5a receptor-antagonist core tetrapeptide.
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13-NOV-1995
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                 (CIBA ) CIBA
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93US-00139913.
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10-OCT-2001;
18-DEC-2001;
The present invention describes a human leukocyte antigen (HLA) (e.g. soluble HLA) ligand database assembled by a process which involves providing a computer system capable of storing HLA data as a database, producing HLA having ligands loaded on it, isolating the loaded ligands from HLA, sequencing the loaded ligands to obtain HLA ligand data, and populating the database with HLA ligand data. Also described: (1) accessing HLA (e.g. sHLA) data stored in a database; and (2) a computer system for HLA (e.g. sHLA) ligand database. The database is populated with HLA ligand sequences, motifs, extended motifs, submotifs, ligands
                                                                                                                                                                                                                                                                                                                                                                     Human leukocyte antigen ligand database assembled by producing HIA having loaded ligands, isolating and sequencing loaded ligands to obtain HIA ligand data and populating database in computer system with ligand data.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C5a receptor antagonists having compsns to treat C5a-mediated c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HILD/)
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MHC ligand;
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                                                                                                                                                                                                                                                                                                           Fig 5; 118pp; English.
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2001US-00974366.
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Pred. No. 1.7e+06;
0; Mismatches 1
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                                                                                                                                                                                                                          Hildebrand WH,
                                                                                                                                                                                                                                                                (HILD/)
(PRIL/)
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10-OCT-2001;
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ligand; major
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HICKMAN H
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                                                                                                                                                                                                                                                                                                    2001US-0270357P.
2001US-00974366.
2001US-00022066.
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histocompatibility
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Pred. No. 1.7e+06;
0; Mismatches 1
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complex; viral; bacterial; tumour.
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The present invention describes a human leukocyte antigen (HLA) (e.g. soluble HLA) ligand database assembled by a process which involves providing a computer system capable of storing HLA data as a database, producing HLA having ligands loaded on it, isolating the loaded ligands from HLA, sequencing the loaded ligands to obtain HLA ligand data, and populating the database with HLA ligand data. Also described: (1) accessing HLA (e.g. sHLA) data stored in a database; and (2) a computer system for HLA (e.g. sHLA) ligand database. The database is populated with HLA ligand sequences, motifs, extended motifs, submotifs, ligands

Human leukocyte antigen ligand database assembled by producing HLA having loaded ligands, isolating and sequencing loaded ligands to obtain HLA ligand data and populating database in computer system with ligand data.

Disclosure; Fig 5; 118pp;

English.

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RESULT 7
ABG96845
AD G96845
AD G96845
AC ABG9
AC ABG9
XX ABG9
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XW Soli
XW Soli
XW Mali
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The invention describes a method of producing soluble human leukocyte antigen (HLA) molecules (SHLA) in cell pharm involving amplifying HLA allelic DNA by PCR using a locus specific primer to produce truncated a PCR product (PI), inserting P1 into mammalian expression vector; electroporating the plasmid into a host cell; inoculating the cell pharm with the host cell such that cell pharm produces sHLA. A multimeric HLA complex (I) is useful for testing functionality of peptide ligands bound
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Soluble human leukocyte antigen; HLA; sHLA; cell pharm; multimeric HLA complex; cytotoxic T lymphocytes; CTL; immune response; viral infection; cancer; autoimmune disease; vaccine development; MHC; major histocompatibility complex; diagnostic development; HLA-BL5 allotype; ligand.
                                                                                                                                                                                                                                                                                                                                   Producing soluble human leukocyte antigen (HLA) in cell pharm useful studies of peptide loading for characterizing human immune responses involves using HLA allelic cDNA or genomic DNA as starting material.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-DEC-2000; 2000US-0256409P
18-DEC-2000; 2000US-0256410P
10-OCT-2001; 2001US-00974366
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                                                                                                                                                                                                                                                                      Disclosure; Page 178; 300pp; English
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1.7e+06;
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Matches 2
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                                                                                           Producing soluble human leustudies of peptide loading involves using HLA allelic
                                                                                                                                                                                                                                                                                             Hildebrand WH,
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18-DEC-2000; 2000US-0256410P
10-OCT-2001; 2001US-00974366
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                                                                                                                                                                                                                            WPI; 2002-698563/75
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                                                                                                                                                                                                                                                                                                                                                      (HILD/) HILDEBRAND W H:
(PRIL/) PRILLIMAN K R.
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2; Conserv
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                                                                                           leukocyte antigen (HLA) in cell pharm useful ing for characterizing human immune responses lic cDNA or genomic DNA as starting material.
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Pred. No. 1.7e
0; Mismatches
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1.7e+06;
1;
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Best Local
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18-DEC-2000; 2000US-0256410P.
10-OCT-2001; 2001US-00974366.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Soluble human leukocyte antigen; HLA; sHLA; cell pharm; multimeric HLA complex; cytotoxic T lymphocytes; CTL; immune responderal infection; cancer; autoimmune disease; vaccine development; major histocompatibility complex; diagnostic development; HLA class I polymorphism; HLA-B15 allotype; ligand.
       WPI; 2002-698563/75
                                                                  Hildebrand WH
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                                                                                                                                 (HILD/)
(PRIL/)
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                                                                                                                              HILDEBRAND W H. PRILLIMAN K R.
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                                                               Prilliman
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Pred. No. 1.7e+06;
0; Mismatches 1;
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levelopment; MHC;
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The invention describes a method of producing soluble human leukocyte antigen (HLA) molecules (sHLA) in cell pharm involving amplifying HLA callelic DNA by PCR using a locus specific primer to produce truncated a CC PCR product (FI), inserting P1 into mammalian expression vector; complex (I) is useful for testing functionality of peptide ligands to complex (I) is useful for testing functionality of peptide ligands bound by at least two soluble HLA molecules. (I) can be tested for its ability companies in humans. (I) is useful for studying T cell responses to pathological conditions such as viral infections and cancer, and for modulating the human immune system to induce tolerance in autoimmune companies produced are useful for studying T cell responses to the individual secreted major histocompatibility complex (MHC) molecules produced are useful for studies of peptide loading (i.e., in concleted MHC molecules, naturally loaded peptides can be cluted from the secreted MHC molecules and functional impact of HLA class I conserved conservation of structural and functional impact of HLA class I conservation of positively or negatively validate the occurrence of identified to positively or negatively validate the occurrence of coverlapping ligands. The truncated version of (MHC) can be produced in mammalian or insect/bacterial cells such that milligram or greater quantities of an individual class I or class II molecule can be obtained. This sequence represents a HLA (human leukocyte antigen) peptide ligand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Producing soluble human leukocyte antigen (HLA) in cell pharm useful studies of peptide loading for characterizing human immune responses involves using HLA allelic cDNA or genomic DNA as starting material.
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Matches 2
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                                    Similarity 2; Conserv
                                    86.7%;
nilarity 66.7%;
Conservative
                                    Score 13; DB 5; Le
Pred. No. 1.7e+06;
0; Mismatches 1;
                                                     Length 4;
                                     Indels
                                    0;
                                    Gaps
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Sequence 4 AA;

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RESULT 10
ABP67652
                                                                                                                                                                                                                                                                                                                                        Human; CD66; CEACAM; cytostatic; antiinflammatory; immunomodulator; antibacterial; virucide; gene therapy; vaccine; neutrophil;
28-FEB-2001; 2001US-0272113P
                                                          27-FEB-2002; 2002WO-US005720
                                                                                                                                                                          WO200268601-A2
                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Human CD66 family modulating peptide SEQ ID NO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-DEC-2002 (first entry)
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                                                                                                                                                                                                                                                                                         immune system; autoimmune disease; cancer; infection; bacterial; virus;
inflammatory disease; transplantation; immunisation.
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(SKUB/) SKUBITZ K M. (SKUB/) SKUBITZ A P N

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RESULT 11
ABP67653
ID ABP67677
XX ABP67653
AD761610-DE
XX Humar
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Matches 2
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New peptide from a surface exposed region of a CD66 family member, useful for modulating the function of CD66 family members, e.g. activation of neutrophils, for treating or diagnosing autoimmune diseases or cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; CD66; CEACAM; cytostatic; antiinflammatory; immunomodulator; antibacterial; virucide; gene therapy; vaccine; neutrophil; immune system; autoimmune disease; cancer; infection; bacterial; virus;
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                                                                                                                                  WPI; 2002-706981/76.
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                                                                                                                                                                                                                                                              (SKUB/) SKUBITZ K M. (SKUB/) SKUBITZ A P N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    inflammatory disease; transplantation;
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Pred. No. 1.7e+06;
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RESULT 12
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ID 84247
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Caspase; apoptosis; GAP SH3 binding protein; HCD2; AOP-1; cancer apoptosis-associated disease; hyperproliferative disease; AIDS degenerative disease; autoimmune disease; Alzheimer's disease;
                                                                                                  Novel apoptosis-associated and/or -modified protein useful as target for the diagnosis, prevention or treatment of apoptosis-associated diseases particularly neurodegenerative diseases.
                                                                                                                                                                                                                                                                        16-JUN-2000; 2000EP-00112813: 16-NOV-2000; 2000EP-00125013:
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Unidentified.
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                                                                                                                                                                                                          Thiede B,
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Pred. No. 1.7e
0; Mismatches
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1.7e+06;
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The invention relates to an apoptosis-associated and/or -modified protein (I) selected from GAP SH3 binding protein, HCD2 and AOP-1 (not defined) or its proteolytic fragments. (I) is useful as target for the diagnosis,

Page 62;

86pp;

English

Å

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RESULT 13
ABJ18638
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Best Local S
Matches 2
The invention comprises peptide antagonists designed to target erbB receptors. The erbB receptor peptide antagonists are useful for preventing tumours and cancers (e.g. gastric cancer, breast cancer and prostate cancer). The present amino acid sequence represents an erbB receptor peptide antagonist of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    prevention or treatment of apoptosis-associated diseases, and in a method for identifying apoptosis modulators. (I) is useful for treatment of hyperproliferative or degenerative diseases including cancers, autoimmune diseases, Alzheimer's disease, viral infections such as acquired immunodeficiency syndrome (AIDS) and vascular diseases such as myocardial infarction. (I) is also useful for treating and/or preventing diseases related to oxidative stress like ischaemic stroke, arthritis, heart failure, Parkinson's disease and amyotrophic lateral sclerosis. (I) is also useful for develop modification-specific diagnostic tools. (I) is also useful for developing protein chips or other solid phase screening devices for high throughput screens, and also as a diagnostic tool to screen for caspase activity, and/or to determine the effectivity of caspase cleavage inhibiting and/or modulating substances. ANU84237-ANU84230 represent caspase cleavage sites of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-APR-2001; 2001US-0282037P
03-AUG-2001; 2001US-0309864P
                                                                                                                                                                          Claim 1; Page 82; 115pp; English
                                                                                                                                                                                                                                          or prostate cancer) in humans
                                                                                                                                                                                                                                                                New peptide antagonists against erbB receptors, useful treating tumors or cancers (e.g. lung adenocarcinomas,
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Greene MI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-APR-2002; 2002WO-US011211.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABJ18638 standard; peptide; 4 AA
                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-075482/07
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (UYPE-) UNIV PENNSYLVANIA
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2; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zhang H,
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Pred. No. 1.7e+06;
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                                                                                                                                                                                                                                                                       for preventing or breast carcinomas
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Sequence 4 AA,

RESULT 15 ADJ81984

ADJ81984 standard;

peptide;

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RESULT 14
ABP59243
ID ABP59
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Best Local S
Matches 2
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Best Local :
                                                                                                 The present invention relates to novel antibodies, which have specificit for activated cell surface receptors (exbB, TWF and immunoglobin gene superfamily IgST) and bind to an epitope (ABB5923-ABB59259). The antibodies are useful for preventing transformation of a normal cell int a tumour cell in a patient who had a tumour with tumour cells that express activated exbB receptors on the cell surface surgically removed or in whom the tumour has entered remission, for treating a patient suffering from a cancer characterised by tumour cells that have pi85 on their cell surfaces, or for treating a patient suffering from an erbB tumour. The antibodies were produced by tumounising a suitable host with peptide (ABP59233-ABP59259), or its reverse sequence
                                                                             Sequence
                                                                                                                                                                                                                                                                                                  Novel antibody for preventing, specificity for activated cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                          Claim 53; Page 88; 114pp; English
                                                                                                                                                                                                                                                                                    an epitope that does not comprise phosphotyrosine residue
                                                                                                                                                                                                                                                                                                                                        WPI; 2003-268117/26.
                                                                                                                                                                                                                                                                                                                                                               Greene MI,
                                                                                                                                                                                                                                                                                                                                                                                                                  03-AUG-2001; 2001US-0309864P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Peptide epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mmunoglobin
1 GXY 3
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                         Similarity 2; Conser
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                                                                             4 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cytostatic;
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                                      86.7%;
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                         ; Score 13; DB; Pred. No. 1.7e
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                               treating, imaging or diagnosing tumor, I surface receptor and binds specifically
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cell surface receptor antibody, SEQ
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                                       DB 6; I
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                                                                                                                            Query Match 86.7
Best Local Similarity 66.7
Matches 2; Conservative
                                                                                                                                                                                                                       The invention relates to novel nucleic acid polymerases from the thermophilic organism Thermus brookianus or mutants having a mutation that decreases 5'-3' exonuclease activity or reduces discrimination against dideoxymucleotide triphosphates. The nucleic acid polymerases, isolated nucleic acids encoding the polymerases, vectors, host cells and kits are useful for DNA synthesis, primer extension, DNA sequencing, reverse transcription, or DNA and RNA amplification procedures. The nucleic acids are also useful for the recombinant expression of the polymerase polymerides. The nucleic acid polymerase and nucleic acids encoding DNA polymerase I activity. This sequence corresponds to a peptide of a polymerase of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nucleic acid polymerase; thermophile; mutation; exonuclease activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Thermus brockianus nucleic acid polymerase peptide fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New Thermus brockianus nucleic acid polymerases, useful for DNA synthesis, primer extension, DNA sequencing, reverse transcription, or DNA and RNA amplification procedures.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Thermus brockianus.
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                                                                                                                                                                                             Sequence 4 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 18; 74pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-505285/47.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rozzelle J, Bolchakova E;
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66.7%;
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Result
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Perfect score:
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       Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Match
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1: /cgn2_6/ptcdata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptcdata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptcdata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptcdata/1/iaa/6B_COMB.pep:*

5: /cgn2_6/ptcdata/1/iaa/FCTUS_COMB.pep:*

6: /cgn2_6/ptcdata/1/iaa/backfiles1.pep:*
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Gapop 10.0 , Gapext 0.5
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12.396 Million cell updates/sec
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Copyright (c) 1993 - 2004 Compugen Ltd
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Sequence 49, Appli
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Query Match Best Local Similar Matches 2; Con	NESULT 1 US-08-798-897-49 Sequence 49, Application US/087 Patent No. 5789201 GENERAL INFORMATION: Genes (TITLE OF INVENTION: Homolog INTATE: DC COUNTRY: USA INTELICATION: HOMOLOG INTATE: DC INTATE: DC COMPUTER: IBM PC Compatil OPERATING SYSTEM: PC-DOSA SOFTWARE: PatentIn Release CURRENT APPLICATION UNDERE: US/08 FILING DATE: February 11, CLASSIFICATION NUMBER: US/08 INTELEPAN: 202-371-2640 INFORMATION FOR SEQ ID NO: 45		29 29 29 30 31 31 31 33 31 33 33 33 33 33
86.7%; Scilarity 66.7%; Pr Conservative 0;	WESSLER, Ork Avenu M: KESSLER, Ork Avenu M: Release MTA: Release MTA: RESALER ORMATION: ATION: AT		77 6 6 2 2 USS 77 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7
red. No. 3.8e+05 Mismatches	g For Bcl-y, OLDSTEIN & FO N.W., Suite OS, Version # ,897 7	ALIGNMENTS	-08-248-538-40 -08-260-199A-31 -08-260-199A-32 -08-690-605-21 -08-580-988A-6 -08-580-988A-6 -08-481-985-21 -08-481-9858-13 -08-932-7878-12 -08-932-7878-12 -09-518-737-6 00320-13 00320-13 00320-13 00320-13 00320-13 00320-13
Length 4; ; 1; Indels	a BC1-2 X P.L.L.C. 600		Sequence
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Patent No.
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TELEX: 139-125
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
LENGTH: 4 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                 Sequence 2, Application US/08463377 Patent No. 5837499
                                                                                                                                                                                                                                                                                                                                          -08-463-377-2
                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patenth Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/463,224
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Schmitz, Albert
APPLICANT: van Heeke, Gino
TITLE OF INVENTION: C5a Receptor Antagonists Having
TITLE OF INVENTION: Substantially No. 5807824Agonist Activity
NUMBER OF SEQUENCES: 67
                                                                                   APPLICANT: van Heeke, Gino
TITLE OF INVENTION: C5a Receptor Antagonists Having
TITLE OF INVENTION: Substantially No. 5837499Agonist Activity
NUMBER OF SEQUENCES: 67
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lerner, David, Littenberg, Krumholz & Mentlik
STREET: 600 South Avenue West
                                                                                                                                                                                                                             APPLICANT: van Oostrum, Jan
APPLICANT: Boyar, William C.
APPLICANT: Galakatos, Nicholas G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: FOLEY, Shawn P.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 908-654-5000
                                                                                                                                                                                              APPLICANT:
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COMPUTER READABLE FORM:
                                             STREET: 600 Sout
CITY: Westfield
STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: U:
ZIP: 07090
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                 COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
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                                  USA
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Schmitz, Albert
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                                                                                                                                                                                                              Schmitz, Albert
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .8e+05;
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US-08-978-523-49
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Best Local Similarity
Thiches 2; Consert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 49,
                                                                                           TELEFAX: 202-371-2540
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: FOLBY, Shawn P.
TELECOMVUNICATION INFORMATION:
TELEPHONE: 908-654-5000
                                                                                                                           NAME: Esmond, Robert W.
REGISTRATION NUMBER: 32,893
REFERENCE/DOCKET NUMBER: 14
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
                                                                                                                                                                                                                                                       CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION UNMEBER: US 08/798,897
FILING DATE: February 11, 1997
                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 53

CORRESPONDENCE ADDRESS:
ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX P.L.L.C.
STREET: 1100 New York Avenue, N.W., Suite 600
                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 4 amino acids
                                                                                                                                                                                                                     CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Guastella, John
TITLE OF INVENTION: Genes
TITLE OF INVENTION: Homolog
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                        STRANDEDNESS:
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J: not relevant
linear
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Pred. No. 3.8e+05;
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; MOLECULE TYPE: peptide US-08-978-523-49

Local Similarity

Pred. No. 3

Length 4;

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                                                                                                                 US-08-310-912A-199
                                                        Matches
                                                                       Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 02110-2904

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30B

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/310,912A

FILING DATE: September 22, 1994

CLASSIFICATION: 536
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                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/227,360
FILING DATE: April 13, 1994
ATTORNEY/AGENT IMPORMATION:
NAME: Lech, Karen F
REGISTRATION NUMBER: 35,238
REGISTRATION NUMBER: 35,238
                                                                                                                                  MOLECULE TYPE:
                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 00
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Kunkel, Barbara N.
APPLICANT: Mindrinos, Michael N.
APPLICANT: Yu, Guo-Liang
TITLE OF INVENTION: RPS2 GENE FAMILY, PRIMERS, PROBES, AND DETECTION
TITLE OF INVENTION: METHODS
MINUSED OF INVENTION: METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
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                                                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                                                        TELEFAX:
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                           1 GXY 3
                                                       Similarity
2; Conserv
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GTY 4
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                                                                                                                                                                                            4 amino acids
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                                                                                                                                                                                                                                                       (617) 542-8906
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Brent, Andrew F.
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                                                                                                                                                             not relevant
                                                                     66.7%;
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                                                                    Score 13; DB 2;
Pred. No. 3.8e+05;
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                                                                                  Length 4;
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APPLICANT: Staskawicz, Brian J.
APPLICANT: Brent, Andrew F.
APPLICANT: Dahlbeck, Douglas
APPLICANT: Katagiri, Fumiaki
APPLICANT: Kunkel, Barbara N.
APPLICANT: Mindrinos, Michael N.
APPLICANT: Yu, Guo-Liang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 41, Application PC/TUS9301669
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Fa
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Vei
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/01669
FILING DATE: 01-MAR-1993
CLASSIFICATION DATA:
APPLICATION NUMBER: US/07/844,852
FILING DATE: 03-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: Wetherell, Jr. Ph.D., John R.
RETERRENCE/DOCKET NUMBER: 31,678
RETERRENCE/DOCKET NUMBER: PD-1636
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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Matches 2; Conserv
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CURRENT FILING DATE: 1999-04-28
EARLIER APPLICATION NUMBER: 08/210,912
EARLIER FILING DATE: 1994-09-22
EARLIER FILING DATE: 1994-09-23
EARLIER FILING DATE: 1994-04-13
NUMBER OF SEQ ID NOS: 208
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ITLE OF INVENTION: RPS2 GENE FAMILY, PRIMERS, ITLE OF INVENTION: DETECTION METHODS ILE REFERENCE: 00786/254002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Trowbridge, Ian S. APPLICANT: Collawn, Jr., Jame APPLICANT: Tainer, John A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                             ZIP: 90067
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: California
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1880 Century Park East, Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Application US/09301085
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Pred. No. 3.8e+05;
                                             , John R.
                                                                                                                                                                                                                              Version
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PCT-US95-04589-199
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Best Local (
Best Local Similarity Matches 2; Conserv
                             Query Match
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TELEFAX: (619) 455-5110
INFORMATION FOR SEQ ID NO:
                                                                                                                              INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 4 amino acids
                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/227
EILING DATE: 13-APR-1994
ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REGISTRATION NUMBER: 0078
TELECHONE: (617) 542-5070
TELEPHONE: (617) 542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 4 amino acids
                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:

MEDIUM TYRE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: FOLDOS/MS-DOS
COPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30B
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Yu, Guo-Liang
TITLE OF INVENTION: RPS2 GENE AND USES THEREOF
NUMBER OF SEQUENCES: 201
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
STREET: 225 Franklin Street Suite 3100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE:
                                                                         MOLECULE TYPE:
                                                                                                                                                                TELEFAX: 100254
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les 2; Conserv
                                                                                TYPE: strandedness: nou
                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: FILING DATE: CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
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(619) 455-5110
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Katagiri, Fumiaki
    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Brent, Andrew
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                                                                       protein
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66.7%;
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                86.7%;
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Pred. No. 3.8e+05;
   Score 13; DB 5;
Pred. No. 3.8e+05;
0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 4;
                              Length 4;
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    Gaps
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Query Match
Best Local Similarity
Matches 2; Conserve
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US-07-962-569A-6
                                                                                                                                                                                                                                                     US-08-292-968-30
                                                                                                                                                                                                                                                                       RESULT 10
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                                                                                                                                                                                                                    Sequence 30, Application US/08292968 Patent No. 5856122
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/07/962,569A
FILING DATE: 19921013
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: PEPPER PH.D.; FREDERICK W.
REGISTRATION NUMBER: 31,286
REFERENCE/DOCKET NUMBER: 920224.01
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: MENON, RAVI S.
APPLICANT: JEFFERS, KATHLEEN F.
APPLICANT: CHANG, YING-FON
APPLICANT: HAM, RICHARD G.
TITLE OF INVENTION: HUMAN K-CAS
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (619) 451-1120
TELEPAX: (619) 451-9628
INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: READ, Randy J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM CC compatible
OPERATING SYSTEM: PC-DDS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 5 amino acids
TYPE: AMINO ACID
TOPOLOGY: linear
MOLECULE TYPE: protein
              TITLE OF INVENTION: MODIFICATION OF PERTUSSIS TOXIN NUMBER OF SEQUENCES: 46
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                   APPLICANT:
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HAM, RICHARD G.
HUMAN K-CASEIN
                                                                               STEIN, Penelope E. CCCKLE, Stephen A. COMEN, Raymond P. LOOSMORE, Sheena KLEIN, Michel H. ARMSTRONG, Glen D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     E: FREDERICK W. PEPPER, PH.D.
11545 W. BERNARDO COURT, STE. 302
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                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                  HAZES, Bart
Sim & McBurney
                                                                                                                                                                                                                                                                                                                                                                                                           86.7%;
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STREET: Suite CITY: Toronto

Suite 701,

330 University Avenue

Ontario

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GENERAL INFORMATION:
APPLICANT: READ, Randy J.
APPLICANT: READ, Penelope E.
APPLICANT: COCKLE, Stephen A.
APPLICANT: COCKLE, Stephen A.
APPLICANT: COCKLE, Stephen A.
APPLICANT: LOOSMORE, Sheena
APPLICANT: KLEIN, Michel H.
APPLICANT: ARMSTRONG, Glen D.
APPLICANT: ARMSTRONG, Glen D.
APPLICANT: HAZES, BEATT
TITLE OF INVENTION: MODIFICATION OF PERTUSSIS TOXIN
NUMBER OF SEQUENCES: 46
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & MCBurney
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:

APPLICATION UNMBER: US 08/110,947

FILING DATE: 24 A UG-1993

PRIOR APPLICATION DATA:

APPLICATION DATA:

APPLICATION UNMBER: US 08/251,121

FILING DATE: 31-MAY-1994

ATTORNEY/AGENT INFORMATION:

NAME: STEWART, Michael I.

REGISTRATION NUMBER: 24/973

REFERENCE/DOCKET NUMBER: 1038-388

TELECOMMUNICATION INFORMATION:

TELEPHONE: (416) 595-1163

INFORMATION FOR SEQ ID NO: 30:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
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ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC COMPS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/292,968
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           -08-292-968-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-292-968-36
COUNTRY: Canada
ZIP: M5G 1R7
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                        STREET: Suite
CITY: Toronto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
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                                                                                                                                                                                      Ontario
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Pred. No. 3.8e+0
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RESULT 12
US-08-618-696-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 66.7%;
Matches 2; Conservative
TELEFAX: 512-474-7577
TELEX: NOT APPLICABLE
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acid residues
                                                                                                 APPLICATION NUMBER: 07/995,269
FILING DATE: 12/21/92
ATTORNEY/AGENT INFORMATION:
NAME: PARKER, DAVID L.
REGISTRATION NUMBER: 32,165
REFERENCE/DOCKET NUMBER: UOAB:002/PAR
TELECOMMUNICATION INFORMATION:
TELEPHONE: 512-320-7200
                                                                                                                                                                                                                                                               COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORDPERECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/618,696
FILING DATE: 20-MAR-1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
ZIP: 77210
COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (416) 595-1155
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 435
PRIOR APPLICATION DATA:
RAPPLICATION NUMBER: US 08,
FILING DATE: 24-AUG-1993
PRIOR APPLICATION DATA:
RAPPLICATION NUMBER: US 08,
RAPPLICATION NUMBER: US 08,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD, WH:
STREET: P.O. BOX 4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: COOPER, Jr., J. ALLEN D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: INHIBITION OF PHAGOCYTES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 31-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Nichael:
REGISTRATION NUMBER: 24;
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: P.O. BUCTON STATE: TEXAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS: single TOPOLOGY: linear
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Pred. No. 3.8e+05;
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Best Local Similarity
Matches 2; Conserv:
                                                                                             RESULT 14
US-08-467-974-30
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Best Local S
Matches 2
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                                                                                                                                                                                                                                                                                                                                     IFLEX: NOT APPLICABLE
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino and Types
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 14, Application US/08618696
Patent No. 5861475
GENERAL INFORMATION:
APPLICANT: COOPER, Jr., J. ALLEN
                                                             Sequence 30, Patent No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:

PRIOR APPLICATION NUMBER: 07/995,269

FILING DATE: 12/21/92

ATTORNEY/AGENT INFORMATION:

NAME: PARKER, DAVID L.

REGISTRATION NUMBER: 32,165

REFERENCE/DOCKET NUMBER: 10AB:002/PAR

TELECOMMUNICATION INFORMATION:

TELEPHONE: 512-320-7200
                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
COMPUTER: IBM PC COMPATIBLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: COOPER, Jr., J. ALLEN D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: INHIBITION OF PHAGOCYTES
NUMBER OF SEQUENCES: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
                                                                                                                                                                                                                                                                                                                  LENGTH: 5 amino acid residues
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: HOUSTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 20
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM:
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                                                                              Application US/08467974
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READ, Randy J.
STEIN, Penelope E.
COCKLE, Stephen A.
                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-MAR-1996
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Pred. No. 3.8e
0; Mismatches
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Pred. No. 3.8e+05;
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3.8e+05;
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US-08-467-974-36
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                                                                                                                                                Patent No. 596:
GENERAL INFORM
APPLICANT:
                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local :
                                                                                                                                                                             Sequence 36, Application US/08467974.
Patent No. 5965385
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 08/251,121
FILING DATE: 31-MAY-1994
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/110,947
FILING DATE: 24-AUG-1993
ATTORNEY/ACENT INFORMATION:
NAME: STEWART, Michael I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-454 MI
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (416) 595-1155
TELEPAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: KLEIN, Mic
APPLICANT: ARMSTROAD
APPLICANT: HAZES, Bas
TITLE OF INVENTION: I
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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LENGTH: 5 amino acids
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APPLICATION NUMBER: 1
FILING DATE: 06-JUN-
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CITY: Toronto
                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/292,968
FILING DATE: 22-AUG-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
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2; Conserva
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                         INVENTION:
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Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OOMEN, Raymond P.
LOOSMORE, Sheena
KLEIN, Michel H.
ARMSTRONG, Glen D.
                                                 STEIN, Penelope E.
COCKLE, Stephen A.
COMEN, Raymond P.
LOOSMORE, Sheena
KLEIN, Michel H.
ARMSTRONG, Glen D.
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                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                          linear
                                                                                                                                                    Randy J
           Bart
: MODIFICATION OF PERTUSSIS TOXIN
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DB 2;

Length 5; Indels

0

Gaps

0

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COUNTRY: CARACA

ZIP: MSG 1R7

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN PC-LOSS/MS-DOS
SOFTWARE: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: IBM PC-DOS/MS-DOS
SOFTWARE: IBM PC-DOS/MS-DOS
SOFTWARE: IBM PC-DOS/MS-DOS
SOFTWARE: APPLICATION DATA:
APPLICATION NUMBER: US/08/467,974
FILING DATE: 06-UN-1995
APPLICATION NUMBER: US/08/467,536
FILING DATE: 06-UN-1995
APPLICATION NUMBER: US/08/292,968
FILING DATE: 22-AUG-1994
PRIOR APPLICATION NUMBER: US/08/291,121
FILING DATE: 31-MAY-1994
PRIOR APPLICATION NUMBER: US/08/251,121
FILING DATE: 24-AUG-1993
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Michaell I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-454 MIS:vg
TELEPRONE: (416) 595-1155
TELEPRONE: GHARACTERISTICS:
LENGTH: 5 amino acids
STRANDEDDESS: single
US-08-467-974-36
Search completed: December 13, 2004, 22:34:51 Job time: 22.4 secs
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Best Local Similarity 66.7%;
Matches 2; Conservative
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ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
CITY: Toronto
                                                                                                                                            1 GXY 3
                                                                                                                                                                                     Score 13; DB 2; Length 5; Pred. No. 3.8e+05; Indels 0; Mismatches 1; Indels
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Page 7

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Database :
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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seq length: 2000000000
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Gapop 10.0 , Gapext 0.5
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1: /cgn2 6/ptodata/2/pubpaa/US07 PUE

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Copyright (c) 1993 - 2004 Compugen Ltd
/ Cgn2_6/ptodata/2/pubpaa/US03_FUBCOMB.pep:*
/ Cgn2_6/ptodata/2/pubpaa/US03E_PUBCOMB.pep:*
/ Cgn2_6/ptodata/2/pubpaa/US03E_PUBCOMB.pep:*
/ Cgn2_6/ptodata/2/pubpaa/US09C_PUBCOMB.pep:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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Sequence 41, Appl	40,	Sequence 36, Appl	Sequence 11, Appl	w.	00 D			34,	199,	Ψ.	Sequence 18, Appl	Sequence 199, App	Description

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-10-375-992-15	-10-264-374-15	9-880-748-3	9-910-552-2	-09-962-445-1	-09-756-283A-9	9-756-283A-5	-09-765-086-15	-10-741-481-3	8-104-32	-10-286-457-1	-10-286-457-12	-09-792-686A-3	-809-517A-1	-10-813-638-1	-10-739-583-1	-10-415-014-7	-10-133-234A-2	-10-133-234A-2	33-234A-	-10-133-234A-1	-10-133-234A-1	-10-435-751-12	-10-435-751-1	-10-306-631-63	-10-394-980-2	-10-284-660-2	0-285-0	-09-858-852A-	-09-858-852A-4	US-09-858-852A-36	
quence 157, Ap	equence 157, 1	equence 3084,	nce 22,	nce 15, App	e 97,	equence 59, App	quence 157, App	equence 35, App	e 326, Ap	equence 189	equence 129, Ap	equence 31,	quence 15, Appl	equence 1446, A	e 14,	equence 732, Ap	equence 21	equence 20, App	quence 19	equence 18, App	equence 17, App	equence 121, Ap	equence 108, Ap	equence 63, App	equence 232, Ap	equence 23, App	equence 23	equence 41	equence 40, App	ence 36	CARCITICA CA

ALIGNMENTS

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APPLICANT: Mindrinos, Michael N.
APPLICANT: Yu, Guo-Liang
ITILE OF INVENTION: RES2 GENE FAMILY, PRIMERS, PROBES, AND
ITILE OF INVENTION: DETECTION METHODS
ITILE OF INVENTION: DETECTION METHODS
ITILE OF INVENTION: DETECTION METHODS
CURRENT FILLING DATE: 2001-05-29
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/301,085
PRIOR FILING DATE: EARLIER FILING DATE: 1999-04-28
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 08/310,912
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 08/310,912
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 08/327,360
PRIOR APPLICATION NUMBER: EARLIER FILING DATE: 1994-04-13
NUMBER OF SEQ ID NOS: 208
SOFTWARE: FRSTSCQ for Windows Version 4.0
                                                                                                       ; LENGTH, 4
; TYPE: PRT
; ORGANISM: Arabidopsis thaliana
US-09-867-852-199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 1
US-09-867-852-199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Ausubel, I
APPLICANT: Staskawid
APPLICANT: Brent, Au
        Query Match
Best Local Similarity 66.
Matches 2; Conservative
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APPLICANT:
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Patent No. US20020147324A1
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Katagiri, Fumiaki
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icz, Brian
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86.7%; Score 13; DB 9; Le
66.7%; Pred. No. 1.4e+06;
tive 0; Mismatches 1;
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                                                                                US-10-613-472-199
                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/613,472
CURRENT FILING DATE: 2003-07-02
PRIOR APPLICATION NUMBER: US 09/667,852
PRIOR PLING DATE: 2001-05-29
PRIOR PELICATION NUMBER: US 09/301,085
PRIOR FILING DATE: 1999-04-28
PRIOR PELICATION NUMBER: US 08/310,912
PRIOR PELICATION NUMBER: US 08/310,912
PRIOR PELICATION NUMBER: US 08/227,360
PRIOR APPLICATION NUMBER: US 08/227,360
PRIOR FILING DATE: 1994-09-22
PRIOR PILING DATE: 1994-04-13
NUMBER: US 08/227,360
PRIOR FILING DATE: 1994-04-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn version 3.2
SEQ ID NO 18
LENGTH: 4
                                                                                                                                                              NUMBER OF SEQ ID NOS: 214
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 199, Application US/10613472 Publication No. US20040088756A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
Matches 2; Conserv
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Best Local Similarity
Matches 2; Conserv
                                       Query Match
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CURRENT FILING DATE: 2002-11-22
PRIOR APPLICATION NUMBER: 60/334,434
PRIOR FILING DATE: 2001-11-30
NUMBER OF SEQ ID NOS: 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Ausubel, Frederick M.
APPLICANT: Staskawicz, Brian J.
APPLICANT: Brent, Andrew F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: THERMUS BROCKIANUS NUCLEIC ACID POLYMERASES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: ROZZELLE, James
APPLICANT: BOLCHAKOVA, Elena
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: 4768US
                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: RPS2 GENE F
TITLE OF INVENTION: DETECTION
FILE REFERENCE: 00786/254004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Thermus brockianus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PPLICANT:
                                                                                                    ORGANISM: Arabidopsis thaliana
                                                                                                                                             ENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mindrinos, Michael N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kunkel, Barbara N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Katagiri,
86.7%;
ilarity 66.7%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Guo-Liang
TON: RPS2 GENE FAMILY, PRIMERS, PROBES,
TON: DETECTION METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fumiaki
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Pred. No. 1.4e
0; Mismatches
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Score 13; DB 15; Length 4; Pred. No. 1.4e+06; 0; Mismatches 1; Indels
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Sequence 34. Application US/09792686A

Publication No. US20030068645A1

GENERAL INFORMATION:
APPLICANT: Kopecky, Joing J.
APPLICANT: Kopecky, Stephen L.
APPLICANT: Weyand, Cornelia M.
TITLE OF INVENTION: METHODS AND MATERIALS FOR EVALUATING TITLE OF INVENTION: CARDIOVASCULAR CONDITIONS
FILE REFERENCE: 07039-250001
CURRENT APPLICATION NUMBER: US/09/792,686A
CURRENT ETLING DATE: 2001-02-23
NUMBER OF SEQ ID NOS: 37
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PRIOR APPLICATION NUMBER: US 09/867,852
PRIOR APPLICATION NUMBER: US 09/867,852
PRIOR APPLICATION NUMBER: US 09/867,852
PRIOR APPLICATION NUMBER: US 09/301,085
PRIOR FILING DATE: 1999-04-28
PRIOR APPLICATION NUMBER: US 08/310,912
PRIOR FILING DATE: 1994-09-22
PRIOR FILING DATE: 1994-09-22
PRIOR APPLICATION NUMBER: US 08/227,360
PRIOR FILING DATE: 1994-04-13
NUMBER OF SEQ ID NOS: 214
SOFTWARE: PASISEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    В
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                                                                      ; TYPE: PRT; ORGANISM: Homo sapiens US-09-792-686A-34
                                                                                                                                                                                                                                                                                                                                                                                                       US-09-792-686A-34
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Publication No. US20040172673A1
                                                                                                                              NUMBER OF SEQ ID NOS: 37
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 34
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Best Local
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                                 Query Match
Best Local Similarity 66.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
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                                                                                                                            ENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SALMERON,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BAKER, BARBARA
ELLIS, JEFFREY
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                 86.7%;
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Pred. No. 1.
Score 13; DB 10; Length 5; Pred: No. 1.4e+06; O; Mismatches 1; Indels
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1.4e+06;
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CURRENT APPLICATION NUMBER: US/10/344,779

CURRENT FILING DATE: 2003-02-17

PRIOR APPLICATION NUMBER: PCT/US01/26161

PRIOR FILING DATE: 2001-08-21

PRIOR APPLICATION NUMBER: US 60/226,524

PRIOR FILING DATE: 2000-08-21

PRIOR APPLICATION NUMBER: US 60/230,639

PRIOR FILING DATE: 2000-09-07

NUMBER OF SEQ ID NOS: 18

SOFTWARE: PastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 143, Application US/10376121A
Publication No. US20030216544A1
GENERAL INFORMATION:
APPLICANT: Harley, John
TITLE OF INVENTION: METHODS AND REAGENTS
AUTOANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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APPLICANT: TORNE
APPLICANT: TRUNE
APPLICANT: WATTE
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Publication No. US20030211106A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: TRUNEH, ALEMSBOED
APPLICANT: WATTAM, TREVOR A.
TITLE OF INVENTION: ANTI-RANK LIGAND MONOCLONAL ANTIBODIES
TITLE OF INVENTION: USEFUL IN TREATMENT OF RANK LIGAND MEDIATED DISORDERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
                                                                               COUNTRY: USA
ZIP: 30309-3400
COMPUTER READABLE FORM:
MEDLIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/10/376,121A
FILING DATE: 27-Mar-2003
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ID NO 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Patrea I. Pabst
STREET: Suite 2000, 1201 West Peachtree Street, N.E.
CITY: Atlanta
                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 GXY 3
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2; Conserv
APPLICATION NUMBER: 07/867,819
FILING DATE: April 13, 1992
APPLICATION NUMBER: 07/648,205
FILING DATE: January 31, 1991
                                                                                                                                                                                                                                                                                                                                      STATE: GA
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Pred. No. 1
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RESULT 9 US-10-394-471B-3

Sequence 3, Application US/10394471B Publication No. US20040185047A1 GENERAL INFORMATION:

Jill; Carton, Jill; Scallon, Bernard J.

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Sequence 8, Application US/10738809
; Publication No. US20040171117A1
; GENERAL INFORMATION:
APPLICANT: SWEET, RAYMOND W.
APPLICANT: TRUNEH, ALEMSEGED
APPLICANT: WATTAM, MARK A.
APPLICANT: WATTAM, TREVOR A.
TITLE OF INVENTION: USEFUL IN TREATMENT OF RANK LIGAND MONOCLONAL ANTIBODIES
TITLE OF INVENTION: USEFUL IN TREATMENT OF RANK LIGAND MEDIAT
CURRENT APPLICATION NUMBER: US/10/738,809
; CURRENT APPLICATION NUMBER: US/10/7344,779
PRIOR APPLICATION NUMBER: PCT/US01/26161
; PRIOR APPLICATION NUMBER: PCT/US01/26161
; PRIOR APPLICATION NUMBER: US 60/226,524
PRIOR APPLICATION NUMBER: US 60/226,524
PRIOR APPLICATION NUMBER: US 60/230,639
; PRIOR FILING DATE: 2000-09-07
; NUMBER OF SEQ ID NOS: 18
; SOFTWARE: FASTSEQ for Windows Version 4.0
; SEQ ID NO 8
; LENGTH: 5
; TYPE: PRT
ORGANISM: Mus musculus
US-10-738-809-8
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US-10-738-809-8
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                                                                           Query Match
Best Local Similarity
Matches 2; Conserv
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Best Local Similarity
Matches 2; Conser
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TELEPAX: (404)-817-8588
INFORMATION FOR SEQ ID NO: 143:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 07/472,947
FILING DATE: January 31, 1990
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 143:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Pabst, Patrea L.
REGISTRATION NUMBER: 31,21
REFERENCE/DOCKET NUMBER: C
TELECOMMUNICATION INFORMATION:
                                     1 GXY 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 5 amino acids
                                                                                  Conservative
                                                                                                  86.7%;
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                                                                               0; Mismatches
                                                                                                Score 13; DB 16; Length 5; Pred. No. 1.4e+06;
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Pred. No. 1.4e+06;
0; Mismatches 1; Indels
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Query Match
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; ORGANISM: Homo sapiens
US-10-394-471B-3
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Best Local Similarity
Matches 2; Conserva
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US-10-394-471B-11
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Sequence 36, Application US/09990762
Patent No. US20020119498A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                     RESULT 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-10-394-471B-11
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PRIOR FILING DATE: 2002-03-26
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PATENTIN VET 3.1
SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 11, Application US/10394471B Publication No. US20040185047A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Giles-Komax, Jill; Carton, Jill; Scallon, Be TITLE OF INVENTION: ANTI-TNF ANTIBODIES, COMPOSITIONS, FILE REFERENCE: CENC202
CURRENT APPLICATION NUMBER: US/10/394,471B
CURRENT FILING DATE: 2003-03-21
PRIOR APPLICATION NUMBER: 60/367,903
PRIOR PILING DATE: 2002-03-26
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PatentIn Ver 3.1
SOFTWARE: P
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CURRENT APPLICATION NUMBER: US/10/394,471B
CURRENT FILING DATE: 2003-03-21
                                 APPLICANT: JOUNG, J. KEITH
APPLICANT: MILLER, JEFFREY
APPLICANT: MILLER, JEFFREY
APPLICANT: PABO, CARL O,
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INTERACTION TRAP ASSAYS
FILE REPERENCE: MTV-030.02 (20021-3002)
CURRENT FILING INTE: 2001-11-14
PRIOR APPLICATION NUMBER: 09/858,852
PRIOR APPLICATION NUMBER: 09/858,852
PRIOR APPLICATION NUMBER: 60/204,509
PRIOR FILING DATE: 2001-05-16
PRIOR FILING DATE: 2000-05-16
NUMBER OF SEQ ID NOS: 97
NUMBER OF SEQ ID NOS: 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: ANTI-THE ANTIBODIES, COMPOSITIONS, METHODS AND USES FILE REFERENCE: CENO202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ENGTH:
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Pred. No. 1.4e+06;
0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 13; DB 16;
Pred. No. 1.4e+06;
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S, METHODS AND USES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Description of Artificial Sequence: Zinc finger; OTHER INFORMATION: recognition sequence US-09-990-762-40
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US-09-990-762-41
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 40, Application US/09990762
Patent No. US20020119498A1
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Patent No. US20020119498A1
GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INTERACTION TRAP ASSAYS FILE REFERENCE: MTV-030.02 (20021-3002)
CURRENT APPLICATION NUMBER: US/09/990,762
CURRENT FILING DATE: 2001-11-14
PRIOR APPLICATION NUMBER: 09/858,852
PRIOR PILING DATE: 2001-05-16
PRIOR APPLICATION NUMBER: 09/858,852
PRIOR PILING DATE: 2001-05-16
PRIOR PILING DATE: 2000-05-16
                                                      SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: JOUNG, J. KEIT APPLICANT: MILLER, JEFFR APPLICANT: PABO, CARL O.
                                                                                        APPLICANT: PABO, CARL O.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INTERACTION TRAP ASSAYS
FILE REFERENCE: METHODS AND COMPOSITIONS FOR INTERACTION TRAP ASSAYS
CURRENT APPLICATION NUMBER: US/09/990,762
CURRENT FILING DATE: 2001-11-14
PRIOR APPLICATION NUMBER: 09/858,852
PRIOR FILING DATE: 2001-05-16
PRIOR PPLICATION NUMBER: 60/204,509
PRIOR FILING DATE: 2001-05-16
PRIOR PPLICATION NUMBER: 60/204,509
PRIOR FILING DATE: 2000-05-16
NUMBER OF SEQ ID NOS: 97
                                                                                                                                                                                                                                                                      APPLICANT: JOUNG, J. KEITH APPLICANT: MILLER, JEFFREY APPLICANT: PABO, CARL O.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 6
TYPE: PRT
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ORGANISM: Artificial Sequence
                                     ENGTH: 6
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2; Conserv
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66.7%;
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Pred. No. 1.4e
0; Mismatches
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Pred. No.
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; FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Zinc finger; OTHER INFORMATION: recognition sequence
US-09-990-762-41

Length 6;

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; LENGTH: 6
; TYPE: PRT
TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Zinc finger
; OTHER INFORMATION: recognition sequence
US-09-858-852A-36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: FUKUI, YASUHISA
APPLICANT: NAGATA, SATOSHI
APPLICANT: SHIRAI, RYUICHI
APPLICANT: SHIRAI, RYUICHI
ITILE OF INVENTION: MONOCLONAL ANTIBODY RECOGNIZING
TITLE OF INVENTION: PHOSPHATIDYLINOSITOL-3,4-DIPHOSPHATE
FILE REPERENCES: 1965/49618
FULRENT APPLICATION NUMBER: US/09/518,737
CURRENT APPLICATION NUMBER: US/09/518,737
CURRENT FILING DATE: 1909-03
PRIOR APPLICATION NUMBER: JP 1999-250209
PRIOR FILING DATE: 1999-09-03
PRIOR FILING DATE: 1999-09-03
PRIOR FILING DATE: 1999-09-03
PRIOR FILING DATE: 1999-09-03
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US-09-858-852A-36
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; ORGANISM: Mus musculus
US-09-518-737-6
                                                                                                                                                APPLICANT: JOUNG, J. KEITH
APPLICANT: MILLER, JEFFREY
APPLICANT: MILLER, JEFFREY
APPLICANT: PABO, CARL O.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INTERACTION TRAP ASSAYS
FILE REFERENCE: MTV-030.01 (20021-3001)
CURRENT APPLICATION NUMBER: US/09/858,852A
CURRENT FILING DATE: 2001-05-16
PRIOR APPLICATION NUMBER: 60/204,509
PRIOR APPLICATION NUMBER: 60/204,509
PRIOR FILING DATE: 2000-05-16
NUMBER OF SEQ ID NOS: 91
SOFTWARE: PATENTIN VEY: 2.1
SEQ ID NO 36
                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 36, Application US/09858852A Publication No. US20030044787A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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Publication No. US20030008321A1
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Pred. No. 1.4e+06;
Mismatches 1; Indels
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Pred. No. 1.4e+06;
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Query Match

86.7%;

Score 13;

DB 10;

Length 6;

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                                    Best Local Similarity Matches 2; Conserv
                  1 GXY 3
GAY 5
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                                    66.7%; Pred. No. 1.4e+06; tive 0; Mismatches 1;
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Search completed: December 13, 2004, 23:02:39 Job time: 69.6 secs

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Result
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Maximum
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Maximum Match 100%
Listing first 45 summaries
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seq length: 2000000000
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Match
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  GenCore version (c) 1993 - 2004
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336850
G41950
PT0214
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unidentified 6.3/4
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ALIGNMENTS

RESULT 1

S36850

Ig heavy chain V region - mouse
C;Species: Mus musculus (house mouse)
C;Date: 06-Jan-1994 #sequence_revision 01-Dec-1995 #text_change 05-Nov-1999
C;Accession: S36850
R;Jacob, J; Kelsoe, G.
R;Jacob, J; Kelsoe, G.
Submitted to the EMBL Data Library, July 1992
submitted to the EMBL Data Library, July 1992
A;Description: In situ studies on the primary immune response to (4-hydroxy-3-nitrophen A;Accession: S36850
A;Accession: S36850
A;Accession: S36850
A;Besidues: preliminary
A;Molecule type: nucleic acid
A;Residues: 1-9 </AC>
A;Cross-references: EMBL:X67387; NID:g50113; PIDN:CAA47799.1; PID:e51594; PID:g1333871
C;Keywords: heterotetramer; immunoglobulin

Similarity 2; Conservat GXY 3 GSY 5	C; Detected Museum Minute Musics (C) Peter (3. Feb. 1994 #text_change 07-May-1999 C; Date: 03-Feb.1994 #text_change 07-May-1999 R; Whetsell, M.; Mosley, R.L.; Whetsell, L.; Schaefer, F.V.; Miller, K.S.; Klein, J.R. Mol. (Call. Biol. 11, 5902-5909, 1991 Mol. (Call. Biol. Biol. 11, 5902-5909, 1991 Mol. (Call. Bio	Qy 1 GXY 3 Db 6 GSY 8 RESULT 2 RESULT 2 G41946 T-cell receptor gamma chain (2t.23) - mouse (fragment)	Query Match 86.7%; Score 13; DB 2; Length 9; Best Local Similarity 66.7%; Pred. No. 2.8e+05; Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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RESULT 3

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A; Molecule t
A; Residues:
C; Keywords:
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C;Date: 30
C;Accessio
R;Yamada,
J. Exp. Me
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J. Exp. Med. 173, 1091-1097, 1991
A;Title: T cell receptor V gene usage of islet beta cell-reactive A;Reference number: PT0209; MUID:91217621; PMID:1902501
A;Accession: PT0214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G. J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT0250
RESULT
S47391
                                                                                                                                                                                                                                                                                                                                                             RESULT
A33099
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A;Experimental source: B lyr
C;Keywords: heterotetramer;
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A; Residues:
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submitted to the Protein Se
A;Reference number: A33098
A;Accession: A33099
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Matches
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;Species: Homo sapiens (man)
;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
;Accession: PT0250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -cell receptor beta chain V-J region (4-1-L.6) - mouse (fragment);Species: Mus musculus (house mouse);Date: 31-Dec-1991 #sequence_revision 31-Dec-1991 #text_change 30-May-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;Molecule type: DNA
;Residues: 1-11 <YAM>
;Experimental source: B lymphocyte
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;Molecule type: prote
;Residues: 1-12 <NIC:
                                                                                                                                                                                                                                                                                                 Accession: A33099
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;Date: 24-Aug-1990 #sequence_revision
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Best Local
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;Residues: 1-11 <NAK>
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Pred. No. 8.8e+02;
0; Mismatches 1
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Pred. No.
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submitted to the EMBL Data Library, August 1994
A;Description: Human HIA-A0201 restricted recognition
A;Reference number: $47355
A;Accession: $47391
A;Status: preliminary
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T-cell antigen receptor VJ junction beta chain - human
C;Species: Homo sapiens (man)
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change
C;Accession: S47395
                                                                                                                              A;Molecule type: DNA
A;Residues: 1-12 <LEV>
A;Experimental source: bone
C;Keywords: immunoglobulin
                                                                                                                                                                                                A;Title: Molecular characterization of transgene-induced A;Reference number: PH1580; MUID:93301609; PMID:8315387 A;Accession: PH1611
                                                                                                                                                                                                                                                                                            Ig H chain V-D-J region (wild-type clone 341) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 02-Jun-1994 #sequence_revision 02-Jun-1994 #text_change 17-Mar-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Cross-references:
C;Keywords: T-cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      submitted to the EMBL Data Library, August 1994
A;Description: Human HLA-A0201 restricted recognition
A;Reference number: 847355
A;Accession: 847395
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                                                                                                                                                                                                                                                R;Levinson, D.A.; Campos-Torres, J.; Leder, J. Exp. Med. 178, 317-329, 1993
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A; Residues: 1-12 < LEH>
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A; Residues: 1-12 <LEH>
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Matches 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cross-references: EMBL:Z35712; NID:g527519; PIDN:CAA84781.1;
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Pred. No. 8.8e
0; Mismatches
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R,Lehner, P.J.
submitted to the EMBL Data Library; August 1994
A;Description: Human HLA-A0201 restricted recognition
A;Reference number: 847355
A;Accession: $47368
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A;Reference number: PH0746; MUID:92078846; PMID:1836010
A;Accession: PH0802
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PH0802
                                        A; Molecule type: mRNA
A; Residues: 1-13 < LEH>
                                                                                A;Status: preliminary
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allelic exclusion and antigen-specific repertoire.
A;Reference number: PH0746; MUID:92078846; PMID:1836010
A;Recession: PH0790
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J. Exp. Med. 174, 1371-1383, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   T-cell receptor alpha chain (QB7.3:2 V-alpha-4.3) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Abate: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PH0790
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A;Experimental source: T lymphocyte
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A; Residues: 1-12 < CAS>
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                                                                                                                                                                                                                          -cell antigen receptor VJ junction beta chain -
,Species: Homo sapiens (man)
,Date: 06-Jan-1995 #sequence_revision 06-Jan-199
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Best Local
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;Residues: 1-12 <CAS>
Cross-references: EMBL:Z35693; NID:g527477; PIDN:CAA84762.1; PID:g527478; Keywords: T-cell receptor
                                                                                                                                                                                                              Accession: S47368
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Experimental source: T lympho
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Pred. No. 8.8e
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Pred. No. 8.8e+02;
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C;Accession: PQ0700
R;Komateu, S.; Kajiwara, H.; Hirano, H.
Theor. Appl. Genet. 86, 935-942, 1993
A;Title: A rice protein library; a data-file
A;Reference number: PQ0696
                                                                    unidentified 6.3/40K protein [imported] - rice (fragment) C;Species: Oryza sativa (rice) C;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text (C;Accession: P00700
                                                                                                                                       RESULT
PQ0700
                                                                                                                                                                                                                                                                                                                                                                                          R:Casanova, J.L.; Romero, P.; Widmann, C.; Kourilsky, P.; Maryanski, J.L.
J. Exp. Med. 174, 1371-1383, 1991
A;Title: T cell receptor genes in a series of class I major histocompatibility complex-allelic exclusion and antigen-specific repertoire.
A;Reference number: BH0746; MUID:92078846; PMID:1836010
A;Accession: PH0788
A;Molecule type: mRNA
A;Residues: 1-13 cCAS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 T-cell antigen receptor VJ junction beta chain - human C,Species: Homo sapiens (man) C,Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #te: C,Accession: S47383
R,Lehner, P.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT
S47383
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A;Experimental source: T lympho
C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text
C;Accession: PH0788
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT
PH0788
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Molecule typė: mRNA
A;Residues: 1-13 <LEH>
A;Cross-references: EMBL:Z35709; NID:g527513; PIDN:CAA84778.1;
C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                submitted to the EMBL Data Library, August 1994
A;Description: Human HLA-A0201 restricted recognition of
A;Reference number: S47355
A;Accession: S47383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Status: preliminary
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66.7%;
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Pred. No. 9.5e.
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RESULT 15
A59018
MUC1 enhancer binding protein 70K chain MUC1EBP-70 - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 08-May-1998 #sequence_revision 08-May-1998 #text_change 08-May-1998
C;Accession: A59018
R;Abe, M; Smith, C.J.; Larson, C.J.
submitted to the Protein Sequence Database, May 1998
A;Description: Involvement of "Ku-like" proteins in the transcription of MUC1/DF3, a bre
A;Reference number: A59018
A;Accession: A59018
A;Mccession: A59018
A;Mccession: A59018
A;Mccession: A59018
A;Mccession: Sequence Database, May 1998
A;Mccession: A59018
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A;Molecule type: protein
A;Residues: 1-13 <KOM>
A;Cross-references: UNIPROT:Q7M1U1
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Search completed: December 13, 2004, 22:32:58 Job time : 16.2 secs
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 si
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1: uniprot_sprot:*
2: uniprot_trembl:*
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Gapop 10.0 ,
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15
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Q47893
Q47893
PPH2 PRUSE
Q7LGD2
Q6JDK5
Q9SMV3
Q6LC19
Q62645
Q911VQ4
Q911VQ4
Q912H6
Q812H6
Q812H6
Q813B14
AAB92363
Q9GJW1
Q9GJW1
Q9GJW1
Q9GJW1
Q9GJW1
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Q9GJW3
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Q9UC46
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     Aac39714 homo sapi
Aab92363 homo sapi
Q9gjw1 lagenorhync
Q9gjw2 lagenorhync
Q9gjw3 lagenorhync
Q61d78 clostridium
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Q6jdk5
Q9smv3
Q9sc19
Q61c19
Q62645
Q71vq4
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Q7y0rl
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Aao63472 homo
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mo sapi
 RESULT 1
UP06_MOUSE
ID UP06_MOUSE
AC P38644;
DT 01-OCT-1994
DT 05-JUL-2004
DT 05-JUL-2004
DE Unknown prott
OS Mus musculus
OC Musmanalia; Bu
OC Mammalia; Bu
OC Mammalia; Bu
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OX NCBI TAXID=11
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RN MEDLINE=9500
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RX MEDLINE=9500
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CC TISGUELAR
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Q9QWT0;
01-MAY-2000 (7
01-MAY-2000 (7
01-DEC-2001 (7
Proteinase 3
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STRAIN-Sv129 D3;

Lutz P.G., Moog-Lutz C., Houzel-Charavel A., Cayre Y.E.

Submitted (JUN-1998) to the EMBL/GenBank/DDBJ databases

EMBL; AJ007030; CAA07429.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE.

SEQUENCE.

SEQUENCE.

SEQUENCE.

MEDLINE=95009907; PubMed=7523108;

Merrick B.A., Patterson R.M., Wichter L.L., He C., Selkirk J.K.;

"Separation and sequencing of familiar and novel murine proteins us

preparative two-dimensional gel electrophoresis.";

Electrophoresis 15:735-745(1994).

-I- MISCELLANEOUS: On the 2D-gel the determined pI of this unknown

protein is: 5.2, its MW is: 50 kDa.

Direct protein sequencing.

NON TER

8 AA; 817 MW; A35DD878676B05B1 CRC64;
                                                                                       NCBI_TaxID=10090;
[1]
                                                                                                                    Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-OCT-1994 (Rel. 30, Created)
01-OCT-1994 (Rel. 30, Last sequence update)
05-UUL-2004 (Rel. 44, Last annotation update)
Unknown protein from 2D-PAGE of fibroblasts (
                                                                                                                                                    Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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(TrEMBLrel. 19,
(Myeloblastin)
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                                                                                                                     Chordata;
Rodentia;
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66.7%;
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Last annotation (Fragment).
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AAB20893
Q9N6I3
Q9GNL6
Q47895
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Q9F6J8
PYRR P1
Q25281
Q9GNL7
Q9S8H8
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Pred. No. 1.8e+06
0; Mismatches
                                                                                                                Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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                                 Cayre Y.E.;
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Q9r4f9 aeromonas.
Q9r501 bacillus su
Q9prx1 struthio ca
Aab20893 clostridi
Q9n6i3 toxoplasma
Q9gn16 drosophila
Q9r4f8 aeromonas.
Q9f6j8 yersinia pe
p37362 pyrrhocoris
Q25261 leishmania
Q9s8h8 brassica na
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Gaps

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Result

Score

Minimum Maximum

BB BB

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Database

Title: Perfect

score:

protein

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Scoring table:

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01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
05-UUL-2004 (TrEMBLrel. 27, Last annotation update)
Type II hair-specific keratin (Type II hair keratin)
Name=KRTHB6;
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NON TER 9
SEQUENCE 9 AA;
                                                                                                                                                                                                                                                                                                                                                         Bairwa N.K., Bamezai R.;
Submitted (DEC-2002) to the EMB
EMBL; AY037552; AAK68688.1; -.
EMBL; AY203963; AAO63472.1; -.
GO; GO:0005882; C:intermediate
                    NON TER
SEQUENCE
                                               SEQUENCE FROM N.A.

Bairwa N.K., Bamezai R.;

Bairwa N.K., Bamezai R.;

"Study of promoter polymorphism

Submitted (DEC-2002) to the EMBI

EMBL; AY203963; AA063472.1; -.
                                                                                                                                      02-MAR-2004 (TrEMBLrel. 27, Creat 02-MAR-2004 (TrEMBLrel. 27, Last 02-MAR-2004 (TrEMBLrel. 27, Last 12-MAR-2004 (TrEMBLrel. 27, Last Type II hair keratin (Fragment).
                                                                                                                                                                                                                                                                                                                                         Keratin.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
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                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A. Bame:
                                                                                                                                                                                                                                                                                                                                                                                                                                 Bairwa N.K., Bamezai
Submitted (MAY-2001)
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                                      Keratin.
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0; Mismatches
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                                                          nism of type II hair keratin gene hHb6.";
EMBL/GenBank/DDBJ databases.
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Pred. No.
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SEQUENCE:
SEQUENCE:
MEDLINE=96326114; PubMed=8703476;
MEDLINE=96326114; Culbreth R.R.;
Copper J.A.Jr., Culbreth R.R.;
Copper J.A.Jr., Culbreth R.R.;
Characterization of a neutrophil inhibitor peptide harvested human bronchial lavage: homology to infiluenza A nucleoprotein. human bronchial lavage: homology 15:207-215(1996).
Am. J. Respir. Cell Mol. Biol. 15:207-215(1996).
GO; GO:0005576; C:extracellular; NAS.
GO; GO:0030236; P:anti-infiammatory response; NAS.
GO; GO:0030236; P:anti-infiammatory response; NAS.
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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01.MAY-2000 (TrEMBLrel. 13, Last sequence update)
01.JUN-2003 (TrEMBLrel. 24, Last annotation update)
Neutrophil inhibitor peptide, NIP=FOLYMORPHONUCLEAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; AY328516; AAP94210.1; NON_TER 1 1
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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01-OCT-2003 (TrEMBLrel. 25, Created)
01-OCT-2003 (TrEMBLrel. 25, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Liu Y., He X., Xu L., Ze Submitted (JUN-2003) to
                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
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Best Local Similarity

Matches 2; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eigenheer R.A., Wiehart U.M., MILLER Bigenheer R.A., Wiehart U.M., MILLER BIGENHEER R.A., Wiehart U.M., MILLER BIGENHEER B.A., "Isolation, identification and localization antidiuretic peptide.", Peptides 24:27-34(2003).

-I- FUNCTION: Inhibitor of fluid secretion in the second messenger. May function is second messenger.
                                                                                                                                                                                               Cell wall.
                                                                                                                                                                                                                                                                                                                                                                         STRAIN-cv. PETIT HAVANA,
Blee K.A., Bonham V.A., Mitchell G.P., R
Wojtaszek P., Bolwell G.P.,
"Proteomic study of secondary cell wall
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Viridiplantae; Streptophyta; Embryo
Spermatophyta; Magnoliophyta; eudicotyledons;
lamiids; Solanales; Solanaceae; Nicotiana.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             P82432;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=4097; [1]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nicotiana tabacum (Common tobacco)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-JUN-2000 (TrEMBLrel. 14, Created)
01-JUN-2000 (TrEMBLrel. 14, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
32 kDa cell wall protein (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Direct protein sequencing; Hormone; Neuropeptide. SEQUENCE 13 AA; 1562 MW; 0240A4504B8A632B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=22465067; PubMed=12576082;
MEDLINE=22465067, Wiehart U.M., Nicolson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADFB_TENMO
P83109;
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                                                                                                                                                                                                                                                                                                                               Planta 0:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                     tobacco culture."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antidiuretic factor B (ADFb).
Tenebrio molitor (Yellow mealworm)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CGMP as second messenger. May function as an antidiuretic SUBCELLULAR LOCATION: Secreted.
TISSUE SPECIFICITY: Brain: Appears to be mainly expressed pairs of bilaterally symmetrical cells in the protocerebrumASS SPECTROMETRY: MW=1560.33; METHOD=MALDI; RANGE=1-13;
                                                                                                                                                                                                                                            TISSUE SPECIFICITY: XYLENGO: 0005618; C:cell wall;
                                                                                                                                                                                                                                                                          SUBCELLULAR LOCATION: CELL WALL TISSUE SPECIFICITY: XYLEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NOTE=Ref.1.
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3 (Rel. 42, Last so
/4 (Rel. 44, Last so
-- factor B (ADFb)
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                                                                  Conservative
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66.7%;
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J.3e+03;
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3.3e+03;
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PIR; PQ0700; PQ0700.
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SEQUENCE 13 AA; 12:
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01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Unidentified 6.3/40% protein (Fragment).
Oryza sativa (Rice).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Trach
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
NCBI TaxID-4530;
                                                                                                                                                                                                                                                                                                                    Ligase.
NON TER
SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                               Wachter A., Steininger H., Rausch T., Bogs C
Submitted (MAY-2003) to the EMBL/GenBank/DDE
EMBL, AJ564376; CAD91995.1; -.
GO; GO:0004357; F:glutamate-cysteine ligase
GO; GO:0016874; F:ligase activity; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eneurosids II; Brassicales; Brassicaceae; Brassica.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-OCT-2003 (TrEMBLrel. 25, Last sequence update)
01-WAR-2004 (TrEMBLrel. 26, Last annotation update)
Gamma-glutamylcysteine synthetase (EC 6.3.2.2) (Fragment)
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07YOR1; PRELIMINARY;
01-OCT-2003 (TrEMBLrel. 25,
01-OCT-2003 (TrEMBLrel. 25,
01-MAR-2004 (TrEMBLrel. 26,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Brassica juncea (Leaf mustard) (Indian mustard)
Eukaryota; Viridiplantae; Streptophyta; Embryop
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Komatsu S., Kajiwara H., Hirano H.;
"A rice protein library; a data-file
two-dimensional electrophoresis";
Theor. Appl. Genet. 86:935-942(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Name=gsh1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=3707;
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e EMBL/GenBank/DDBJ
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Pred. No.
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P80620;
01-OCT-1996 (Rel. 34, Created)
01-OCT-1996 (Rel. 34, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Unknown protein from 2D-PAGE of etiolated coleoptile
(Fragment).
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta;

Spermatophyta; Magnoliophyta;

Spermatophyta; Daniel Control Co
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Touzet P., Riccardi F., Morin C., Damerval C.,
Pernollet J.-C., Zivy M., de Vienne D.;
"The maize two dimensional gel protein databass
genome analysis program.";
Theore analysis program.";
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Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
PACCAD clade; Panicoideae; Andropogoneae; Zea.
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"Prunus serotina amygdalin hydrolase and prunasin hydrolase.";
Plant Physiol. 100:282-290(1992).
-!- CATALYTIC ACTIVITY: (R)-prunasin + H(2)O = mandelonitrile +
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Li C.P., Swain
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MISCELLANEOUS: On the 2D-gel the MISCELLANEOUS; 5.9, its MW is: 34.6
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MEDLINE-86233345; PubMed=3086870;

CONLEY P.B., Lemaux P.G., Lomax T.L., Grossman A.R.;

Conley e.B., Lemaux P.G., Lomax T.L., Grossman A.R.;

"Genes encoding major light-harvesting polypeptides are
the genome of the cyanobacterium Fremyella diplosiphon.

Proc. Natl. Acad. Sci. U.S.A. 83:3924-3928(1986).

EMBL; M13216; AAAA24871.1; -.
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GO: 00.00422; F:beta-glucosidase activity; IEA.

GO: 00.00422; F:beta-glucosidase activity; IEA.
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01-MAY-2000 (TrEMELrel. 13, Last sequence update)
01-JUN-2003 (TrEMELrel. 24, Last annotation update)
Beta-D-glucoside glucohydrolase (EC 3.2.1.21) (Fragment)
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Q47893

PRELIMINARY;

PRT;

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AC Q47893;
DT 01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
DT 01-NOV-1996 (TrEMBLrel. 08, Last annotation update)
DT 01-NOV-1998 (TrEMBLrel. 08, Last annotation update)
DE Beta-phycocyanin (Fragment).
OS Fremyella diplosiphon (Galotthrix PCC 7601).
OS Fremyella diplosiphon (Galotthrix PCC 7601).
OC Bacteria; Cyanobacteria; Nostocales; Microchaetaceae; Fremyella.
OX NCBI_TAXID=1197;
RN [1]
RP SEQUENCE FROM N.A.
RY MEDLINE-86233345; PubMed=3086870;
RA COnley P.B., Lemaux P.G., Lomax T.L., Grossman A.R.;
CONLEY P.B., Lemaux P.G., Lomax T.L., Grossman A.R.;
Conley P.B., Lemaux P.G., Lomax T.L., Grossman A.R.;
RT "Genes encoding major light-harvesting polypeptides are clustered on "RT" the genome of the cyanobacterium Fremyella diplosiphon.";
RT proc. Natl. Acad. Sci. U.S.A. 83:3924-3928(1986).
Proc. Natl. Acad. Sci. U.S.A. 83:3924-3928(1986).
FT NON TER 1
SEQUENCE 15 AA; 1528 MW; 7FF2F65518F493D4 CRC64;
Query Match
Best Local Similarity 66.7%; Pred. No. 9.4e+03;
Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;
Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;
DB 4 GSY 6

Search completed: December 13, 2004, 22:31:36

Job time: 85.8 secs
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Title:
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1: geneseqp1980s:*
2: geneseqp1990s:*
3: geneseqp2000s:*
4: geneseqp2001s:*
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Gapop 10.0 ,
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geneseqp2004s:*
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(c) 1993 - 2004
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    AAP61656

AAP761656

AAP761333

AAR55333

AAR55333

AAR55713

AAP655716

AAB9569697

AAB967657

AAB967657

AAB967657

AAB967653

AAU84247

AAB75924

AAU98858

AAU98858

AAU98858

AAU42068

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Compugen
Aap61656 Sequence
Aap71313 Peptide C
Aar55333 Inhibitor
Aar75499 C5a recep
Abp55796 B15 class
Abp55796 B15 class
Abp55796 B15 class
Abp96697 Human leu
Abg96697 Human CD6
Aau84247 Caspase c
Abj18638 ErbB rece
Abj18638 ErbB rece
Abj5984 Thermus b
Adl9854 Human leu
Adl9854 Human leu
Adl9858 Human leu
Adl9858 Human leu
Adl9858 Human leu
Adl9858 Antibacte
Adk42068 Antibacte
Adk42068 Antibacte
Aar43432 Ro/SSA ep
Aar56398 Synthetic
Aar72555 Pertussis
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being printed,
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25-MAR-2003 03-OCT-2002 21-AUG-1991

(revised)
(revised)
(first entry)

AAP61656;

AAP61656 standard; protein; 4 AA.

Sequence of peptide which inhibits cyclic-nucleotide kinase activity and mammalian cell growth.

independent

protein

45	44	43	42	41	40	39	38	37	36	ω σ	34	i Lu	32	ω L	30	29	28	27	26
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ALIGNMENTS

RESULT 1
AAP61656
ID AAP6
XX AAP6
XX AAP6
AC AAP6
XX AAP6
AC AAP6
XX AAP6
AC AAP6
XX AAP6
AC AAP6
XX Call
XX C WPI; 1986-118872/18. Kettner CA, Racker E; 16-NOV-1983; 16-NOV-1983; Misc-difference US4582821-A. Misc-difference Synthetic. Cell growth inhibitor; tumour cell (DUPO) DU PONT DE NEMOURS & CO 15-APR-1986 83US-00552255 83US-00552255 Location/Qualifiers /label= Tyr-CH2Cl /label= Carbobenzoxy-Phe H growth inhibitor.

The cpds. of the invention inhibit protein phosphorylation. The inventorial process for inhibiting the growth of tumour calls in a medium which comprises contacting the cells with a cpd. of formula (Ahp61654-P61661) or a physiologically acceptable salt. (Updated on 03-OCT-2002 add missing OS field.) (Updated on 25-MAR-2003 to correct PA field.)

The inventors

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Claim 1;

Col 4; 9pp; English.

Inhibition of tumour cell growth - using peptide and aminoacid halo: $methyl\ ketone(s)$.

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RESULT 2
AAP713
AC AAP7
AX 25-M
DT 25-M
DT 15-M
XX Pept
XX Picc
XX Synt
XX VS4M
XX Picc
XX Synt
XX CDU
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Peptide component of cpd. for treating picornavirus infections
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15-MAY-1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-AUG-1984;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Picornaviridae; poliovirus; rhinovirus; antiviral agent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAP71313 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 4 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This peptide is useful as part of a peptide/halo-methyl ketone cpd., for treating picornavirus, egpolio- or rhinovirus infections. It inhibits the processing of picornavirus capsid proteins by virus encoded proteases. See AAP71301-12. See also US4652552. (Updated on 25-MAR-2003 to correct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 4; 10pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Treating picorna-virus infection with peptide halo:methyl ketone cpds. esp. for treating polio virus and rhino virus infections.
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                                                                          25-MAR-2003
30-JAN-1995
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                                                                                                                                                                                               AAR55333 standard; peptide; 4 AA.
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                            Inhibitor of
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2; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                Similarity 66. 2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GAY 4
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                                                                                                                                                                                                                                                                                                                        GAY 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Korant BD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (revised)
(first entry)
                         tyrosine kinase
                                                                        (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein; 4 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                            86.7%;
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Pred. No. 1.7e+06;
0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                        0
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 13; DB 1; 1
Pred. No. 1.7e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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RABSULT 4
AAR754
AAR754
AAR76
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AAR77
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DT 25-M
DT 25-M
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C5a
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PN W099
PN W099
PN 15-C
XX
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PD 15-C
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XX
PD 15-C
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PD 16-N
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PR 16-N
PR 06-I
PR XX
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AAR76
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The sequence is that of a protein tyrosine kinase inhibitor contg. D. Tyr or terrafluoro-Tyr residues. The peptide can be used to inhibit TK-mediated signal transduction and is useful for controlling proliferative diseases, e.g. cancer, psoriasis and to treat viral, inflammatory, allergic and cardiovascular diseases. See also AARS5302-39. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New peptide inhibitors of protein tyrosine kinase - contain D-Tyr or tetrafluoro-Tyr residue, for treating proliferative diseases, viral infection, inflammation etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-NOV-1992;
27-OCT-1993;
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Modified-site
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                                                                                                                                                                                                                                                                                                                          C5a; C5a receptor-antagonist; antiinflammatory.
                                                                                                                                                                                                                                                                                                                                                                               C5a receptor-antagonist core tetrapeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR75499 standard; peptide; 4 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 7; Page 36; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1994-183419/22.
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                                                                                                                                                                                                                                                                               Homo sapiens
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13-NOV-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR75499;
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                                                                             06-DEC-1993;
                                                                                                                             16-NOV-1994;
                                                                                                                                                                             15-JUN-1995
                             (CIBA ) CIBA GEIGY AG
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2; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                  (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      92US-00973136
93US-00139913
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                                                                               93US-00162591
                                                                                                                             94WO-IB000359
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              86.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      8
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Pred. No. 1.7e+06;
0; Mismatches 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (D/L) "
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Boyar WC,

Galakatos NG,

Peppard

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RESULT 5
ABP55713
ID ABP5
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Best Local S
Matches 2
The present invention describes a human leukocyte antigen (HLA) (e.g. soluble HLA) ligand database assembled by a process which involves providing a computer system capable of storing HLA data as a database, producing HLA having ligands loaded on it, isolating the loaded ligands from HLA, sequencing the loaded ligands to obtain HLA ligand data, and populating the database with HLA ligand data. Also described: (1) accessing HLA (e.g. sHLA) data stored in a database, and (2) a computer system for HLA (e.g. sHLA) ligand database. The database is populated with HLA ligand sequences, motifs, extended motifs, submotifs, ligands
                                                                                                                                                                                                                                                                                                       loaded
ligand
                                                                                                                                                                                                                                                      Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hildebrand WH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-FEB-2001; 2001US-0270357P.
10-OCT-2001; 2001US-00974366.
18-DEC-2001; 2001US-00022066.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (HILD/)
(PRIL/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 B15 class I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABP55713 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Polypeptide receptor antagonists to C5a contain an essential tetrapeptide (AAR75498 or AAR75499) or core tripeptide (DGA) displays C5a blocking activity. (Updated on 25-MAR-2003 to compared to the contains of the contains 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C5a receptor antagonists having compsns. to treat C5a-mediated c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-FEB-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1995-224319/29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ligand;
ligand;
                                                                                                                                                                                                                                                                                          leukocyte antigen ligand database assembled by producing HLA having d ligands, isolating and sequencing loaded ligands to obtain HLA data and populating database in computer system with ligand data.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N
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2; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HILDEBRAND W H.
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                                                                                                                                                                                                                                                      Fig 5; 118pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         human
major
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sHLA molecule ligand related peptide
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histocompatibility
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
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Pred. No. 1.7e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hickman
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               predictive algorithm; database;
complex; viral; bacterial; tumour.
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1.7e+06;
1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                correct
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                                        unique to infected cells, tumour specific ligands, as well as a collection of current and future developed HLA ligand sequences. The database which contains endogenously bound and loaded ligands facilitates searching of viral, bacterial, tumour or human protein sequences for ligands likely to bind a particular HLA class I or class II protein. The database allows the user to screen an unknown peptide sequence for potential matches with sHLA ligand discrete sequences or SHLA ligand motifs of sequences. Due to the completeness and concentration of sHLA obtained to date, better sequencing data of numerous endogenously loaded HLA ligands is found in the sHLA ligand database, and by comparison of such ligands to each other and to the genomic sequence, better motifs are also found in the sHLA ligand database. ABP55912 represent amino acid sequences used in the exemplification of the present invention
Sequence 4 AA;
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Query Match
Best Local S
Matches 2
N
             1 GXY 3
                            . Similarity 2; Conserv
                                    86.7%;
66.7%;
                                    Score 13;
Pred. No.
                             Mismatches
                                   DB 5; I
1.7e+06;
                                         Length
                            Indels
                            0
                           Gaps
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B15 class I
                                                                          ABP55796;
                                                                                                         ABP55796 standard; peptide; 4 AA.
                                        25-FEB-2003
sHLA molecule ligand related peptide #90.
                                   (first entry)
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HLA ligand; MHC ligand; human leukocyte antigen; major histocompatibility predictive algorithm; complex; viral; bacte bacterial;

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Synthetic
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RESULT 6
ABP55796
ID ABP5
XX ABP5
XX ABP5
XX ABP5
XX HLA
XW HLA
XW MHC
XX Synt
XX WO20
XX Synt
XX WO20
XX Synt
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10-OCT-2001;
18-DEC-2001;
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2001US-00974366.
2001US-00022066.
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Hildebrand WH, Prilliman ₩, Hickman

HICKMAN H

U K

WPI; 2002-732755/79.

Human leukocyte antigen ligand database assembled by producing loaded ligands, isolating and sequencing loaded ligands to obtaligand data and populating database in computer system with ligand ucing HLA having D obtain HLA Ch ligand data.

Disclosure; Fig 5; 118pp; English

The present invention describes a human leukocyte antigen (HLA) (e.g. soluble HLA) ligand database assembled by a process which involves providing a computer system capable of storing HLA data as a database, producing HLA having ligands loaded on it, isolating the loaded ligands from HLA, sequencing the loaded ligands to obtain HLA ligand data, and populating the database with HLA ligand data. Also described: (1) accessing HLA (e.g. sHLA) data stored in a database; and (2) a computer system for HLA (e.g. sHLA) ligand database. The database is populated with HLA ligand sequences, motifs, extended motifs, submotifs, ligands

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ARBSULT 7
ABG96845
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XX ABG9
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XX Solu
KW Solu
KW Mult
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10-OCT-2001;
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The invention describes a method of producing soluble human leukocyte antiqen (HIA) molecules (sHLA) in cell pharm involving amplifying HLA allelic DNA by PCR using a locus specific primer to produce truncated a PCR product (PI), inserting P1 into mammalian expression vector; electroporating the plasmid into a host cell; inocularing the cell pharm produces sHLA. A multimeric HLA complex (I) is useful for testing functionality of peptide ligands bound
                                                                                                                                                                                                                                                                                                               Producing soluble human leukocyte antigen (HLA) in cell pharm useful studies of peptide loading for characterizing human immune responses involves using HLA allelic cDNA or genomic DNA as starting material.
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Pred. No. 1.7e+06;
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to serve as ligands for cytotoxic T lymphocytes (CTLs) and induce immune cresponses in humans. (I) is useful for studying T cell responses to compatibility the human immune system to induce tolerance in autoimmune compatibility complex (MHC) diseases. The individual secreted major histocompatibility complex (MHC) diseases. The individual secreted major histocompatibility complex (MHC) complex complex (MHC) are represented MHC molecules, naturally loaded peptides can be eluted from the secreted MHC molecules, naturally loaded peptides can be eluted from the complex complex compared to fix control and to the development of diagnostics. With the secreted MHC molecules and functional impact of HLA class I complex complex complex compared to fix class I standard peptide pools extracted from series of distinct yet compared class I HLA-BIS allotypes; compare the different ligands and hence contentify potentially shared elements; and characterise the elements comparished to positively or negatively validate the occurrence of coverlapping ligands. The truncated version of (MHC) can be produced in control contents of an individual class I or class II molecule can be obtained. This sequence represents a HLA (human leukocyte antigen) peptide ligand can be control contents.
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18-DEC-2000;
10-OCT-2001;
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                                         producing soluble human leukocyte antigen (HLA) in cell pharm useful studies of peptide loading for characterizing human immune responses involves using HLA allelic cDNA or genomic DNA as starting material.
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2001US-00974366
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Disclosure; Page 183;

Producing soluble human leukocyte antigen (HLA) in cell pharm useful studies of peptide loading for characterizing human immune responses involves using HLA allelic cDNA or genomic DNA as starting material.

for

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RESULT 9
ABG96697
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CC electroporating the plasmid into a host cell; inoculating the cell pharm coulet such that cell pharm produces sHLA. A multimeric HLA complex (I) is useful for testing functionality of peptide ligands bound by at least two soluble HLA molecules. (I) can be tested for its ability complex (I) is useful for studying T cell responses to pathological conditions such as viral infections and cancer, and for complex complex (I) is useful for studying T cell responses to pathological conditions such as viral infections and cancer, and for comparison the human immune system to induce tolerance in autoimmune complex (MHC) complex (MHC) and to the development of diagnostics. With the secreted MHC molecules, naturally loaded peptides can be eluted from the complex complex (MHC) molecule and characterised. The secreted MHC molecules are also useful to generate ligands and hence ligand maps from the peptide pools extracted from series of distinct yet identify potentially shared elements; and characterise the elements contentially shared elements; and characterise the elements of dentified to positively or negatively validate the occurrence of coverlapping ligands. The truncated version of (MHC) can be produced in mammalian or insect/bacterial cells such that milligram or greater content content is sequence represents a HLA (human leukocyte antigen) peptide ligand
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Matches 2
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                                                                                                                                                   18-DEC-2000; 2000US-0256409P.
18-DEC-2000; 2000US-0256410P.
10-OCT-2001; 2001US-00974366.
WPI; 2002-698563/75
                                           Hildebrand WH,
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PRILLIMAN K F
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llarity 66.7%;
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Pred. No. 1.7e+06;
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The invention describes a method of producing soluble human leukocyte cantigen (HLA) molecules (BHLA) in cell pharm involving amplifying HLA called in DNA by PCR using a locus specific primer to produce truncated a PCR product (PI), inserting P1 into mammalian expression vector; CC electroporating the plasmid into a host cell; inoculating the cell pharm cc with the host cell such that cell pharm produces sHLA. A multimeric HLA complex (I) is useful for testing functionality of peptide ligands bound by at least two soluble HLA molecules. (I) can be tested for its ability to serve as ligands for cytotoxic T lymphocytes (CTLs) and induce immune cresponses in humans. (I) is useful for studying T cell responses to pathological conditions such as viral infections and cancer, and for compatiting the human immune system to induce tolerance in autoimmune conseases. The individual secreted major histocompatibility complex (MHC) consequence produced are useful for studies of peptide loading (i.e., in consequence) and to the development of diagnostics. With the corrected MHC molecules, naturally loaded peptides can be eluted from the medicules produced are useful for studies of peptide loading (i.e., in consequence) and characterised. The secreted MHC molecules allow the secreted MHC molecules allow the conjugation of structural and functional impact of HLA class I conjugate the different ligands and hence conjugated class I HLA-BIS allotypes; compare the different ligand maps to identify potentially shared elements; and characterise the elements conjugatively or negatively validate the occurrence of coverlapping ligands. The truncated version of (MHC) can be produced in mammalian or insect/pacterial cells such that milijaram or greater conjugatively or insect/pacterial cells such that milijaram or greater in the produced in the conjugatively conjugated by the conjugatively produced in the conjugatively conjugated by the conjugated conjugated by the conjugated conjugated by the conjugated conjugated by the conjugated conju
Sequence 4 AA;
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Best Local
                       Similarity
2; Conserv
          GXY 3
                       Conservative
                            86.7%;
                     ; Score 13; DB; Pred. No. 1.7e
                  DB 5;
1.7e+06;
1;
                                  Length
                      Indels
                                  4:
                     0
                     Gaps
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RESULT 10
ABP67652
ID ABP67
Human CD66 family modulating peptide SEQ ID
                                                                                 10-DEC-2002 (first entry)
                                                                                           ABP67652;
                                                                                                  ABP67652 standard; peptide; 4
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                                                                          523.
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Human; CD66; CEACAM; cytostatic; antiinflammatory; immunomodulator; antibacterial; virucide; gene therapy; vaccine; neutrophil; immune system; autoimmune disease; cancer; infection; bacterial; virus; inflammatory disease; transplantation; immunisation.
28-FEB-2001; 2001US-0272113P
                                                                    27-FEB-2002; 2002WO-US005720
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(SKUB/) SKUBITZ K M. (SKUB/) SKUBITZ A P N.

and as a vaccine

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ABP67653
ID ABP67653
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     therapies and
New peptide from a surface exposed region of a CD66 family member, useful for modulating the function of CD66 family members, e.g. activation of neutrophils, for treating or diagnosing autoimmune diseases or cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human CD66 family modulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABP67653;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABP67653 standard; peptide; 4 AA
                                                                                                      WPI; 2002-706981/76
                                                                                                                                                        Skubitz KM,
                                                                                                                                                                                                                                                                           28-FEB-2001; 2001US-0272113P
                                                                                                                                                                                                                                                                                                                                                                            06-SEP-2002
                                                                                                                                                                                                                                                                                                                             27-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                           WO200268601-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity nes 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 a vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 GXY 3
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                                                                                                                                                                                                    SKUBITZ K M.
SKUBITZ A P N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 30; 96pp; English
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                                                                                                                                                        Skubitz APN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                2002WO-US005720
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          for immunisation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 86.7%;
66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               transplantation; immunisation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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Pred. No. 1.7e+06;
0; Mismatches 1
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The invention relates to an isolated peptide (I) from a surface exposed CC region of a CD66 family member (ABP67130-ABP67990). The peptide CC modulates: (a) activation of neutrophils; (b) activation or inhibition of CC T-cells, B-cells, NK cells, LAK cells, dendritic cells or other immune CC ells, including epithelial cells; (d) homotypic and/or heterotypic CC cells, including epithelial cells; (d) homotypic and/or heterotypic CC dathesion among CD66 family members; and (e) adhesion of CD66 family cC members to other ligands. The peptide is useful in modulating the CC function of CD66 family members and/or their ligands, such as activation cf ineutrophils and activation, inhibition, proliferation and/or construction of the immune cells. The peptides may also be used in treating or diagnosing autoimmune diseases, cancer, infections (e.g. bacterial or viral) or inflammatory diseases, in transplantation covered the construction of the immunication diseases, in transplantation covered the construction of the immunication diseases, in transplantation covered the construction of the immunication diseases, in transplantation covered the construction of the immunication diseases, in transplantation covered the construction of the immunication diseases, in transplantation covered the construction in the construction of the construction of the immunication diseases, in transplantation covered the construction of the construction of the immunication diseases.
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Best Local S
Matches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Caspase; apoptosis; GAP SH3 binding protein; HCD2; AOP-1; cancer; apoptosis associated disease; hyperproliferative disease; AID5; degenerative disease; autoimmune disease; Alzheimer's disease; viral infection; acquired immunodeficiency syndrome; vascular disease; myocardial infarction; oxidative stress; ischaemic stroke; arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Caspase cleavage site #11.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAU84247 standard; peptide;
                                                                                                                                                                                                                                                                                                16-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200196873-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         heart failure; Parkinson's disease; amyotrophic lateral sclerosis
                                                                                                                                                                                                                                                            (PLAC ) MAX PLANCK GES FOERDERUNG WISSENSCHAFTEN
                                                                                                                                                                                                                                                                                                                                                        15-JUN-2001; 2001WO-EP006780
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2; Conserv
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                                                                                                                                                                                                                       Thiede B, Machuy N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AA,
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                                                                                                                                                                                                                                                                                                2000EP-00112813.
2000EP-00125013.
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Pred. No. 1.7e
0; Mismatches
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The invention relates to an apoptosis-associated and/or -modified protein (I) selected from GAP SH3 binding protein, HCD2 and AOP-1 (not defined) or its proteolytic fragments. (I) is useful as target for the diagnosis,

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English.

Novel apoptosis-associated and/or -modified protein useful as target for the diagnosis, prevention or treatment of apoptosis-associated diseases particularly neurodegenerative diseases.

2002-098088/13.

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RESULT 13
ABJ18638
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
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The invention comprises peptide antagonists designed to target erbB receptors. The erbB receptor peptide antagonists are useful for preventing tumours and cancers (e.g. gastric cancer, breast cancer and prostate cancer). The present amino acid sequence represents an erbB receptor peptide antagonist of the invention
                                                                                                                                                                                                                                                     New peptide antagonists against erbB receptors, useful treating tumors or cancers (e.g. lung adenocarcinomas, or prostate cancer) in humans,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-APR-2001; 2001US-0282037P. 03-AUG-2001; 2001US-0309864P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ErbB receptor antagonist peptide #21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Greene MI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-OCT-2002.
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                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-075482/07
                                                                                                                                                                                        Claim 1; Page 82; 115pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (UYPE-) UNIV PENNSYLVANIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Murali R,
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 5; I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Berezov
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                                                                                                                                                                                                                                                                                          for preventing or breast carcinomas
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Sequence 4 AA

RESULT 15 ADJ81984

ADJ81984;

ADJ81984 standard; peptide; 4 AA.

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RESULT 14
ABP59243
ID ABP59
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Best Local
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Best Local
                                                          Matches
                                                                                                                                         The present invention relates to novel antibodies, which have specificit for activated cell surface receptors (erbB, TMF and immunoglobin gene superfamily: IGSF) and bind to an epitope (ABP59223-ABP59259). The antibodies are useful for preventing transformation of a normal cell int a tumour cell in a patient who had a tumour with tumour cells that express activated erbB receptors on the cell surface surgically removed or in whom the tumour has entered remission, for treating a patient suffering from a cancer characterised by tumour cells that have pl85 on their cell surfaces, or for treating a patient suffering from an experience of the control of the antibodies were produced by immunising a suitable host with peptide (ABP59223-ABP59259), or its reverse sequence
                                                                                                                                                                                                                                                                                                                                                     Novel antibody for preventing, treating, imaging or diagnosing tumor, k specificity for activated cell surface receptor and binds specifically an epitope that does not comprise phosphotyrosine residue.
                                                                                                                    Sequence 4 AA;
                                                                                                                                                                                                                                                                                                                             Claim 53; Page 88; 114pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-268117/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antibody; cytostatic; vaccine; immunoglobin gene superfamily;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-MAY-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2003012072-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Peptide epitope for generating cell surface receptor antibody, SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABP59243;
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                                                                      86.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Richter M,
                                                     Score 13; DB 6; Le
Pred. No. 1.7e+06;
0; Mismatches 1;
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Pred. No. 1.7e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cell surface receptor; erbB; TNF;
IgSF; epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Murali R;
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                                                           Indels
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                                                                                                                              Query Match 86.7%;
Best Local Similarity 66.7%;
Matches 2; Conservative
                                                                                                                                                                                                                            The invention relates to novel nucleic acid polymerases from the thermophilic organism Thermus brockianus or mutants having a mutation that decreases 5'-3' exonuclease activity or reduces discrimination against dideoxymucleotide triphosphates. The nucleic acid polymerases, isolated nucleic acids encoding the polymerases, vectors, host cells and kits are useful for DNA synthesis, primer extension, DNA sequencing, reverse transcription, or DNA and RNA amplification procedures. The nucleic acids are also useful for the recombinant expression of the polymerase polymerase increases. The nucleic acid polymerases and nucleic acids encoding DNA polymerase I activity. This sequence corresponds to a peptide of a polymerase of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Thermus brockianus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleic acid polymerase; thermophile; mutation; exonuclease activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Thermus brockianus nucleic acid polymerase peptide fragment #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New Thermus brockianus nucleic acid polymerases, useful for DNA synthesis, primer extension, DNA sequencing, reverse transcription, or DNA and RNA amplification procedures.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; SEQ ID NO 18; 74pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rozzelle J, Bolchakova E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-NOV-2001; 2001US-0334434P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-NOV-2002; 2002WO-US037657.
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                                                                                                                                                                                                Sequence 4 AA;
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                                                                                                                                   Score 13; DB 7; Pred. No. 1.7e+06; 0; Mismatches 1
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 g
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Perfect score:
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1: /cgn2 6/ptodata/1,

2: /cgn2 6/ptodata/1,

3: /cgn2 6/ptodata/1,

4: /cgn2 6/ptodata/1,

5: /cgn2 6/ptodata/1,

6: /cgn2 6/ptodata/1,
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length: 2000000000
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Match
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/cgm2_6/ptodata/1/iaa/6A_COMB.pep:*
/cgm2_6/ptodata/1/iaa/6B_COMB.pep:*
/cgm2_6/ptodata/1/iaa/6B_COMB.pep:*
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                   US-08-798-897-49
US-08-433-377-2
US-08-433-377-2
US-08-978-523-49
US-08-978-523-49
US-08-911-085-199
US-08-911-085-199
US-09-910-085-199
US-08-952-968-36
US-08-618-696-3
US-08-618-696-3
US-08-467-974-36
US-08-467-974-36
US-08-467-974-36
US-08-467-974-36
US-08-467-975-36-36
US-08-467-976-36
US-08-082-514-36
US-08-082-514-36
US-08-08-176-620A-21
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Sequence 49, Appli
Sequence 2, Appli
Sequence 2, Appli
Sequence 199, App
Sequence 30, Appli
Sequence 31, Appli
Sequence 31, Appli
Sequence 36, Appli
Sequence 36, Appli
Sequence 37, Appli
Sequence 36, Appli
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Sequence 31, Appli
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                                                                                                                                         STREET: 1100 New York AVenue, ......

CITY: Washington
STATE: DC
COUNTRY: USA
ZIP: 20005

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIAN PC-DOS/MS-DOS
SOFTWARE: PATENTIAN Release #1.0, Version #1..
CURRENT APPLICATION NUMBER: US/08/798,897
FILING DATE: PEDILORY 11, 1997
CLASSIFICATION INFORMATION:
NAME: ESMOND, KOBERT W.
REGISTRATION NUMBER: 32,893
REFERENCE/DOCKET NUMBER: 1483.0140001
TELEPHAN: 202-371-2540
INFORMATION FOR SEQ ID NO: 49:
SEQUENCE CHARACTERISTICS:
LENGTH: 4 amino acids
TYPE: amino acids
TYPE: amino acids
TYPE: amino acids
TYPE: amino acid
STRANNEDNIESS: not relevant
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-798-897-49
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US-08-798-897-49
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Best Local S
Matches 2
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Guastella, John
TITLE OF INVENTION: Genes Coding For Bcl-y, a
TITLE OF INVENTION: Homologue
NUMBER OF SEQUENCES: 53
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
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                                                                                                           l Similarity
2; Conserv
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                                                                                                               Conservative
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US-08-260-199A-32
US-08-260-199A-32
US-08-690-695-21
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US-08-690-695-21
US-08-988-13-23
US-08-988-13-23
US-08-988-818-13-2
US-08-988-818-13-2
US-08-988-818-13-12
US-08-988-818-13-15
US-08-988-818-13-15
US-08-988-818-13-15
US-08-988-818-13-15
US-08-988-818-13-15
US-08-988-818-13-15
US-08-988-818-73-6
US-09-181-997A-19
US-09-193-104-28
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Pred. No. 3.8e
0; Mismatches
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Sequence 31, Appl
Sequence 53, Appl
Sequence 51, Appl
Sequence 16, Appl
Sequence 17, Appl
Sequence 13, Appl
Sequence 13, Appl
Sequence 12, Appl
Sequence 12, Appl
Sequence 12, Appl
Sequence 12, Appl
Sequence 14, Appl
Sequence 16, Appli
Sequence 19, Appl
Sequence 28, Appli
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Minimum : Maximum :

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Patent No. 580782
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Best Local
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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APPLICANT:
APPLICANT:
                                   STREET: 600 CORNER Westfield
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: FOLEY, Shawn P.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 908-654-5000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/463,224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 07090
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: van Heeke, Gino
TITLE OF INVENTION: C5a Receptor Antagonists Having
TITLE OF INVENTION: Substantially No. 5807824Agonist Activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Lerner, David, Littenberg, Krumholz & Mentlik
STREET: 600 South Avenue West
COMPUTER READABLE FORM:
                                                                                                               APPLICANT: van Heeke, Gino
APPLICANT: van Heeke, Gino
TITLE OF INVENTION: C5a Receptor Antagonists Having
TITLE OF INVENTION: Substantially No. 5837499Agonist Activity
TITLE OF SECUENCES: 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
TOPOLOGY: linear
                           COUNTRY:
                                                                          ADDRESSEE: Lerner, David, Litt
STREET: 600 South Avenue West
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                  07090
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Galakatos, Nichol
Schmitz, Albert
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Schmitz, Albert
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                                                                                                  David, Littenberg, Krumholz & Mentlik
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Pred. No. 3.8e+05;
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Patent No. 5883229
GENERAL INFORMATION:
APPLICANT: Guastella, John
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TELEX: 139-125
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 4 aming acids
            TELEFAX: 202-371-2540
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Esmond, Robert W.
REGISTRATION NUMBER: 32,893
REFERENCE/DOCKET NUMBER: 1483
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION UMMBER: US/08/978,523
FILING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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NAME: Foley, Shawn P.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 908-654-5000
                                                                                                                                                                                CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/798,897
FILLING DATE: February 11, 1997
CLASSIFICATION: 424
CLASSIFICATION: 424
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 66.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX P.L.L.C. STREET: 1100 New York Avenue, N.W., Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                    20005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        g
4 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                      USA
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66.7%;
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Pred. No. 3.8e+05;
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TYPE: amino acid STRANDEDNESS: not

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                                                                                                             ; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-310-912A-199
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                                                     Query Match
Best Local Similarity
Matches 2; Conserv
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APPLICANT:
APPLICANT:
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                                                                                                                                                                                                              TELEX: 100254
INFORMATION FOR SEQ ID NO:
                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 4 amino acids
TYPE; amino acid
STRANDEDNESS: not releva
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: 02110-2904
ZIP: 02110-2904
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30B
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/22
FILING DATE: April 13, 19
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                      NAME: Lech, Karen F.
REGISTRATION NUMBER: 35,238
REFERENCE/DOCKET NUMBER: 00.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Yu, Guo-Liang
TITLE OF INVENTION: RPS2 GENE FAMILY, PRIMERS, PROBES, AND DETECTION
TITLE OF INVENTION: METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                               TELEPHONE: (617) 542-8906
                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE: Septem CLASSIFICATION: 536
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5981730
                          GXY 3
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                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mindrinos, Michael N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Brent, Andrew F
                                                                                                                                                                                                                                                                                                                                                                                                   September 22, 1994
                                                                                                                                                     not relevant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Barbara N
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13, 1994
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                                                    Score 13; DB:
Pred. No. 3.8e
0; Mismatches
                                                     <u>,</u>
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Pred. No. 3.8e+05;
0; Mismatches 1
                                              DB 2; __
3.8e+05;
1;
                                                                              Length 4;
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                                                     Indels
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RESULT 7
PCT-US93-01669-41
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; TITLE OF INVENTION: RPS2 GENE FAMILY, PRIMER
; TITLE OF INVENTION: DETECTION METHODS
; FILE REFERENCE: 00786/254002
; CURRENT APPLICATION NUMBER: US/09/301,085
; CURRENT FILING DATE: 1999-04-28
; EARLIER APPLICATION NUMBER: 08/310,912
EARLIER FILING DATE: 1994-09-22
; EARLIER FILING DATE: 1994-04-13
; NUMBER OF SEQ ID NOS: 208
; SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-301-085-199
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Ver
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/01669
FILING DATE: 01-MAR-1993
CLASSIFICATION DATA:
APPLICATION UNMBER: US/07/844,852
FILING DATE: 03-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: Wetherell, JT. Ph.D., John R.
REGISTRATION NUMBER: 31,678
REFERENCE/DOCKET NUMBER: 9D-1636
TELECOMMUNICATION INFORMATION:
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Best Local :
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TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: RECEPTOR INTERNALIZATION SIGNALS NUMBER OF SEQUENCES: 57
                                                                                                                                                                                                                                                                                                                                                                                                   STREET:
CITY: I
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                                                                                                                                                                                                                                                                                                                                             90067
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California
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Katagiri, Fumiaki
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1880 Century Park East, Suite 500
                                                                                                                                                                                                                                                                                                                                                               USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Collawn, Jr.,
Tainer, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Trowbridge, Ian S
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Pred. No. 3.8e+05;
0; Mismatches 1
                                                                                                                                                                                                                                          Version
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PCT-US95-04589-199
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                                                   TOPOLOGY: linear;
MOLECULE TYPE: protein
PCT-US95-04589-199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 199, Applica GENERAL INFORMATION: AUSUDE APPLICANT: AUSUDE
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INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
  Matches
                           Query Match
                                                                                                                                               TELEX: 100254
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                 ZIP: 02110-2904
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30B
CURRENT APPLICATION NUMBER: PCT/US95/04589
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
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                                                                                                                                                                       REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
TELEPAX: (617) 542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                 CLASSIFICATION:
PRIOR APPLICATION NUMBER: 1
APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             UMBER OF SEQUENCES:
                                                                                                                                                                                                                                            FILING DATE: 13-APR-1994
ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PPLICANT: Yu, Guo-Liang
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TELEX: 100254
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                                                                                           STRANDEDNESS:
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              Local
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2; Conserva
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  Conservative
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Mindrinos, Michael N.
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E: peptide
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                                                                                           not relevant
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Fumiaki
               86.7%;
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  0;
  Score 13; DB 5; Length 4; Pred. No. 3.8e+05; O; Mismatches 1; Indels
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Pred. No. 3.
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RESULT 9
US-07-962-569A-6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 6,
                                                                                                                                                                                   Sequence 30, App
Patent No. 58561
                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: MENON, RAVI S.
APPLICANT: MEPERS, KATHLEEN
APPLICANT: CHANG, YING-FON
APPLICANT: HAM, RICHARD G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (619) 451-9628
INFORMATION FOR SEQ ID NO: (
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
ATTORNEY AGENT INFORMATION:
NAME: PEPPER PH.D., FREDERICK W.
NAME: 31,286
REGISTRATION NUMBER: 31,286
REFERENCE/DOCKET NUMBER: 920224.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION: TELEPHONE: (619) 451-1120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
NUMBER OF SEQUENCES: 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
FILING DATE: 199210
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                        Local Similarity
tes 2; Conserv
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ZIP: 92127
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                  INVENTION: MODIFICATION OF PERTUSSIS TOXIN
                                                                                                                                                                                                   Application US/08292968
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USA
                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                KLEIN, Michel H. ARMSTRONG, Glen D.
                                                                                   LOOSMORE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19921013
                                  Bart
                                                                                                                    Penelope E., Stephen A.
                                                                                                 Raymond P.
                                                                                                                                                                                                                                                                                                                                                                            86.7%;
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                                                                                 Sheena .
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Pred. No. 3.8e+05;
0; Mismatches 1
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STREET: Suite 701,

Toronto : Ontario

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS

APPLICATION NUMBER: US/08/292,968 FILING DATE:

CLASSIFICATION: 435

```
STREET: Suite 701, 330 University Avenue
CITY: Toronto
STATE: Ontario
COUNTRY: Canada
ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PStentIn Release: #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/292,968
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                           E: Sim & McBurney
Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                           MODIFICATION OF PERTUSSIS TOXIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 13; DB 2;
Pred. No. 3.8e+05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 5;
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US-08-618-696-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PILING DATE: 24-AUG-1993
PRIOR APPLICATION DATA:
APPLICATION UNMER: US 08/251
PILING DATE: 31-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Michael I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038
REFERENCE/DOCKET NUMBER: 1038
REJECOMUNICATION INFORMATION:
TELEPHONE: (416) 595-115
TELEPHONE: (416) 595-1163
INFORMATION FOR SEQ ID NO: 36:
SEQUENCE CHARACTERISTICS:
FENERAL CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 65...
""" tohes 2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 3, Application US/08618696 Patent No. 5861475 GENERAL INFORMATION:
APPLICANT: COOPER, Jr., J. ALLEI
                                        TELEX: NOT APPLICABLE INFORMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS: LENGTH: 5 amino acid residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: COM-
TITLE OF INVENTION: INH-
NUMBER OF SEQUENCES: 2.1
CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD, WH-
STREET: P.O. BOX 4433
CITY: HOUSTON
                                                                                                                                         NAME: PARKER, DAVID L.
REGISTRATION NUMBER: 32,165
REFERENCE/DOCKET NUMBER: UO:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 512-320-7200
                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/618,696
FILING DATE: 20-MAR-1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/995,269
FILING DATE: 12/21/92
                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 08/110,947
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE.
COUNTRY: UC.
77210
                                                                                                                 TELEPHONE: 512-320-72
TELEFAX: 512-474-7577
        STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 5 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ARNOLD, WHITE & DURKEE O. BOX 4433
      single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Jr., J. ALLEN D. COMPOSITIONS AND METHODS FOR THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INHIBITION OF PHAGOCYTES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US 08/251,121
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                                                                                                                                                                                   UOAB:002/PAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 5;
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RESULT 11 US-08-292-968-36

Sequence 36, App Patent No. 58561 GENERAL INFORMA

Application US/08292968

INFORMATION:

APPLICANT: READ, Randy J.
APPLICANT: STEIN, Penelope E.
APPLICANT: COCKLE, Stephen A.
APPLICANT: COMEN, Raymond P.
APPLICANT: LOOSMORE, Sheena
APPLICANT: KLEIN, Michel H.
APPLICANT: TRUETONYC COLUMN

APPLICANT: OOMEN, RAPPLICANT: LOOSMORE, APPLICANT: KLEIN, KLEIN, APPLICANT: ARMSTROM APPLICANT: HAZES, BUTTLE OF INVENTION:

HAZES, Bart

ARMSTRONG, Glen D.

CORRESPONDENCE ADDRESS:

ADDRESSEE:

UMBER OF SEQUENCES:

밁 Ş

US-08-292-968-30

LENGTH: 5 amino acids
TYPE: amino acid
STRANDEDNESS: single

linear

Query Match
Best Local Similarity
Matches 2; Conserv

Conservative

9

1 GXY 3

PRIOR APPLICATION UNMBER: US 08/110,947
FILING DATE: 24-AUG-1993
PRIOR APPLICATION NUMBER: US 08/251,121
FILING DATE: 31-MAY-1994
APPLICATION NUMBER: US 08/251,121
FILING DATE: 31-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Michael I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-388
TELECOMMUNICATION INFORMATION:
TELEPAN: (416) 595-1155
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 30:
SEQUENCE CHARACTERISTICS:
IFNORTH: 58-1170

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; TOPOLOGY: US-08-618-696-3
                                         US-08-467-974-30
; Sequence 30, Application US/08467974
; Patent No. 5965385
; GENERAL INFORMATION:
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                                                                                                           RESULT 14
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US-08-618-696-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 14, Application US/08618696
Patent No. 5861475
                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 512-474-7577
TELEX: NOT APPLICABLE
INFORMATION FOR SEQ ID NO:
 APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/ALL.

NAME: PARKER, DAVID D.

NAME: PARKER, DAVID D.

REGISTRATION NUMBER: 32,165
REFERENCE/DOCKET NUMBER: UOAB:002/PAR
REFERENCE/DOCKET NUMBER: UOAB:002/PAR
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION: 512-320-7200
TELECOMMUNICATION: 512-320-7200
                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acid residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/95
FILING DATE: 12/21/92
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORDERREECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: FLOPPY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD, WHITE & DURKEE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: FILING DATE: 20-MAI CLASSIFICATION: 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: P.O. I
                                                                                                                                                                                                                                                                                              STRANDEDNESS:
TOPOLOGY: li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION:
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                                                                                                                                                                                                                 2; Conservative
                                                                                                                                                                                   GXY 3
                                                                                                                                                                                                                                                                                                                       amino acid
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READ, Randy J.
STEIN, Penelope E.
COCKLE, Stephen A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COOPER, Jr., J. ALLEN D.
MENTION: COMPOSITIONS AND METHODS
TENTION: INHIBITION OF PHAGOCYTES
                                                                                                                                                                                                                                                                                            linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-MAR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FLOPPY DISK
                                                                                                                                                                                                                                                                                                           single
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                                                                                                                                                                                                                                   86.7%;
66.7%;
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Pred. No. 3.8e+05;
0; Mismatches 1
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Pred. No. 3.8e+05;
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US-08-467-974-36
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                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 2
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                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
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Sequence 36, Application US/08467974 Patent No. 5965385
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 30:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/4
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/2
FILING DATE: 22-AUG-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/2
FILING DATE: 31-MAY-1994
PRIOR APPLICATION DATA: 198/1
PRIOR APPLICATION DATA: 08/1
PRIOR APPLICATION NUMBER: US 08/1
PRIOR APPLICATION NUMBER: US 08/1
APPLICATION NUMBER: APPLICATION: NUMBER: STEWART, Michael I
                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: Canada
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
TELEPHNE: (416) 595-1155
                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 66. 
nes 2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Toronto
STATE: Ontario
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER:
INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OOMEN, Raymond P.
LOOSMORE, Sheena
KLEIN, Michel H.
ARMSTRONG, Glen D.
                                                                                                    STEIN, Penelope E. COCKLE, Stephen A. OOMEN, Raymond P. LOOSMORE, Sheena
                                         HAZES, Bart
                                                        KLEIN, Michel H.
ARMSTRONG, Glen D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         single
                                                                                                                                                                                     Randy J
                                                                              Michel H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MODIFICATION OF PERTUSSIS TOXIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                    86.7%;
  MODIFICATION OF PERTUSSIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US 08/251,121
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Pred. No. 3.8e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Saim & McBurney
STREET: Suite 701, 330 University Avenue
CITY: Toronto
STATE: Ontario
COUNTRY: Canada
ZIP: MSG1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IMP C compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTWMARE: Patentin Release #1.0, Version #1.25
APPLICATION NUMBER: US/08/467,974
FILING DATE:
CLASSIFICATION LOATA:
APPLICATION NUMBER: US/08/467,536
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/292,968
FILING DATE: 02-AUG-1994
PRIOR APPLICATION NUMBER: US 08/292,968
FILING DATE: 31-MAY-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/251,121
FILING DATE: 31-MAY-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/251,121
FILING DATE: 31-MAY-1994
PRIOR APPLICATION NUMBER: US 08/251,121
FILING DATE: 24-AUG-1993
ATTORNEY/ACENT INFORMATION:
NAME: STEWART NUCHBER: 1038-454 MIS:vg
TELEFORM: (416) 595-1163
INFORMATION FOR SEQ ID NO: 36:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
STYPE: amino acids
STEANDEDNESS: single
US-08-467-974-36
Search completed: December 13, 2004, 22:34:51 Job time: 21.4 secs
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                                                                                                                          Ś
                                                                                                                                                           Query Match 86.3
Best Local Similarity 66.3
Matches 2; Conservative
                                                                                                                          1 GXY 3
                                                                                       GTY 4
                                                                                                                                                                                     86.7%;
                                                                                                                                                           Score 13; DB 2; Length 5;
Pred. No. 3.8e+05;
0; Mismatches 1; Indels
                                                                                                                                                             0; Gaps
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Result
No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Minimum DB
Maximum DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Run on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OM protein -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                     and is derived
                                                                                                                                                                                                                                                                                                                                                                                                                                   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed,
                                                                                                                                                                                                                                                                                                            Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    length: 0 length: 2000000000
                                                                                                                                                                                                                                                                                                         Query
Match
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                             Published_Applications
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Copyright (c) 1993 - 2004 Compugen Ltd
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/ogn2_6/ptodata/2/pubpaa/PCT_I
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gn2-6/ptodata/2/pubpaa/US08_NEW_PUB_Pep:*
gn2-6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
gn2-6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
gn2-6/ptodata/2/pubpaa/US09A_PUBCOMB.pep:*
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gn2-6/ptodata/2/pubpaa/US10C_PUBCOMB.pep:*
gn2-6/ptodata/2/pubpaa/US10C_NEW_PUB.pep:*
gn2-6/ptodata/2/pubpaa/US10C_NEW_PUB.pep:*
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_6/ptodata/2/pubpaa/PCTUS_PUBCOMB.pep:*
_6/ptodata/2/pubpaa/US08_NEW_PUB.pen:*
_6/ptodata/2/pubpaa/US08_NEW_PUB.pen:*
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9 US-09-867-852-199
4 US-10-302-817A-18
5 US-10-613-472-199
6 US-10-613-765-199
9 US-99-792-86A-34
4 US-10-376-17A-143
6 US-10-376-17A-143
6 US-10-374-471B-13
6 US-10-394-471B-3
6 US-10-394-471B-11
6 US-10-394-471B-11
7 US-10-394-471B-11
8 US-09-990-762-41
8 US-09-990-762-41
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  Sequence 199, App
Sequence 19, Appl
Sequence 199, App
Sequence 34, Appl
Sequence 34, Appl
Sequence 8, Appli
Sequence 8, Appli
Sequence 3, Appli
Sequence 11, Appl
Sequence 40, Appl
Sequence 40, Appl
Sequence 40, Appl
Sequence 40, Appl
Sequence 41, Appl
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Œ	equence 157,	equence 30	equence 22,	nce 15,	equence 97,	e 59,	equence 157	Sequence 35, Appl	320	189	129	31	15,	equence 14	equence 14,	equence 73:	equence 21	equence 20,	equence 19,	e 18	equence 17	e 12:	equence 10	equence 63	quence 23:	equence 23	equence 23	equence 41	e 40	equence 36	-

ALIGNMENTS

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RESULT 1
US-09-867-852-199
                                                                                                             ; ORGANISM: Arabidopsis thaliana US-09-867-852-199
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                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/867,852.

CURRENT FILING DATE: 2001-05-29

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/301,085

PRIOR FILING DATE: EARLIER FILING DATE: 1999-04-28

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 08/310,912

PRIOR FILING DATE: EARLIER FILING DATE: 1994-09-22

PRIOR FILING DATE: EARLIER FILING DATE: 1994-09-22

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 08/227,360

PRIOR FILING DATE: EARLIER FILING DATE: 1994-04-13

NUMBER OF SEQ ID NOS: 208

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 199
Query Match 86.7%; Score 13; DB 9; Length 4; Best Local Similarity 66.7%; Pred. No. 1.4e+06; Matches 2; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Kunkel, Barbara N.
APPLICANT: Mindrinos, Michael N.
APPLICANT: Yu, Guo-Liang
TITLE OF INVENTION: RPS2 GENE FAMILY, PRIMERS,
TITLE OF INVENTION: DETECTION METHODS
FILE REFERENCE: 00786/254002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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Kunkel, Barbara N.
Mindrinos, Michael N.
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Fumiaki
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                                                                                   ; TYPE: PRT , ORGANISM: Arabidopsis thaliana US-10-613-472-199
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US-10-302-817A-18
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CURRENT APPLICATION NUMBER: US/10/613,472
CURRENT FILING DATE: 2003-07-02
PRIOR APPLICATION NUMBER: US 09/867,852
PRIOR FILING DATE: 2001-05-29
PRIOR APPLICATION NUMBER: US 09/301,085
PRIOR FILING DATE: 1999-04-28
PRIOR APPLICATION NUMBER: US 08/310,912
PRIOR APPLICATION NUMBER: US 08/310,912
PRIOR APPLICATION NUMBER: US 08/227,360
PRIOR FILING DATE: 1994-04-13
NUMBER OF SEQ ID NOS: 214
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                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 199
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Matches 2; Conservative
Best Local Similarity Matches 2; Conserv
                                        Query Match
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APPLICANT: ROZZELLE, James
APPLICANT: BOLCHAKOVA, Elena
TITLE OF INVENTION: THERMUS BROCKIANUS NUCLEIC ACID POLYMERASES
FILE REFERENCE: 4768US
CURRENT APPLICATION NUMBER: US/10/302,817A
CURRENT FILING DATE: 2002-11-22
PRIOR APPLICATION NUMBER: 60/334,434
PRIOR FILING DATE: 2001-11-30
NUMBER OF SEQ ID NOS: 58
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APPLICANT: Staskawicz, Br
APPLICANT: Brent, Andrew
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Katagiri, Fumiaki
Kunkel, Barbara N.
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icz, Brian J.
                      86.7%;
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  Score 13; DB 15; Length 4;
Pred. No. 1.4e+06;
0; Mismatches 1; Indels
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: KCDCKY, JOYG J.
APPLICANT: KCDCKY, Stephen L.
APPLICANT: KCPCKY, Stephen L.
APPLICANT: KCPCKY, Stephen L.
APPLICANT: KCPCKY, Stephen L.
APPLICANT: KCPCKY, Stephen L.
APPLICANT: KCPCANTION: METHODS AND MATERIALS FOR EVALUATING
TITLE OF INVENTION: CARDIOVASCULAR CONDITIONS
FILE REFERENCE: 07039-250001
CURRENT APPLICATION NUMBER: US/09/792,686A
CURRENT APPLICATION NUMBER: US/09/792,686A
CURRENT FILING DATE: 2001-02-23
NUMBER OF SEQ ID NOS: 37
SOSTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 34
LENGTH: 5
TYPE: DET
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CURRENT APPLICATION NUMBER: US/10/613,765
CURRENT FILING DATE: 2003-07-02
CURRENT FILING DATE: 2003-07-02
PRIOR APPLICATION NUMBER: US 09/867,852
PRIOR APPLICATION NUMBER: US 09/301,085
PRIOR APPLICATION NUMBER: US 09/301,085
PRIOR FILING DATE: 1999-04-28
PRIOR FILING DATE: 1999-04-28
PRIOR APPLICATION NUMBER: US 08/310,912
PRIOR APPLICATION NUMBER: US 08/227,360
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US-10-613-765-199
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                                                                                                                             ; TYPE: PRT; ORGANISM: Homo sapiens US-09-792-686A-34
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US-09-792-686A-34
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Query Match
Best Local Similarity bb..
Conservative
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Publication No. US20040172673A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: RPS2 GENE FAMILY, PRIMERS, PROBES, AND TITLE OF INVENTION: DETECTION METHODS FILE REFERENCE: 00786/254005
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TYPE: PRT
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SALMERON, JOHN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MINDRINOS, MICHAEL N
YU, GUO-LIANG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STASKAWICZ, BRIAN
KATAGIRI, FUMIAKI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BAKER, BARBARA
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                                                        86.7%;
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                                   0; Mismatches
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Pred. No. 1.4e+06;
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US-10-376-121A-143
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APPLICANT: Harley, John
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT FILING DATE: 2003-02-17
PRIOR APPLICATION NUMBER: ETT/US01/26161
PRIOR FILING DATE: 2001-08-21
PRIOR APPLICATION NUMBER: US 60/226,524
PRIOR APPLICATION NUMBER: US 60/230,639
PRIOR APPLICATION NUMBER: US 60/230,639
PRIOR APPLICATION NUMBER: US 60/230,639
PRIOR PILING DATE: 2000-09-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 18
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: GP50033
CURRENT APPLICATION NUMBER: US/10/344,779
CURRENT FILING DATE: 2003-02-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NPPLICANT: WATTAM, TREVOR A.
TITLE OF INVENTION: ANTI-RANK LIGAND MONOCLONAL ANTIBODIES
TITLE OF INVENTION: USEFUL IN TREATMENT OF RANK LIGAND MEDIATED DISORDERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BNGTH:
                                                                                                        COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
COPERATING PATENTIA Rélease #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/376,121A
FILING DATE: 27-Mar-2003
CLASSIFICATION: CUNKNOWN>
PRIOR APPLICATION DATA: 07/867,819
APPLICATION NUMBER: 07/867,819
FILING DATE: April 13, 1992
APPLICATION NUMBER: 07/648,205
FILING DATE: January 31, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Patrea L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 GXY 3
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                                                                                                                                                                                                                                                                                                                                                    CITY: Atlanta
STATE: GA
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                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                   STREET: Suite 2000,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INVENTION: METHODS AND REAGENTS FOR DIAGNOSIS OF AUTOANTIBODIES
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1201 West Peachtree Street, N.E.
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Pred. No. 1.4e+06;
0; Mismatches 1; Indels
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US-10-394-471B-3
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                    Sequence 3, Application US/10394471B Publication No. US20040185047A1 GENERAL INFORMATION:
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Best Local Similarity 66.7
Matches 2; Conservative
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APPLICANT: SWEET, F
APPLICANT: TORNETT
                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                 Query Match
Best Local S
APPLICANT: Giles-Komar, Jill; Carton, Jill; Scallon, Bernard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 2000-09-07
                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Mus musculus
                                                                                                                                                                                                                                                          Local Similarity hes 2; Conserv
                                                                                                                                                                                                                   1 GXY 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TORNETTA, MARK A.
                                                                                                                                                                                                                                                                 Conservative
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NUMBER OF SEQ ID NOS: 18
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 8
LENGTH: 5
TYPE: PRT
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Publication No. US20040171117A1
                                                                                                                                                                                                                                                                        APPLICANT: TRUMER, ALEMSEGED
APPLICANT: WATTAM, TREVOR A.

ITLE OF INVENTION: ANTI-RANK LIGAND MONOCLONAL ANTIBODIES
TITLE OF INVENTION: USSFUL IN TREATMENT OF RANK LIGAND MEDIATED DISORDERS
FILE REFERENCE: GD5003
CURRENT APPLICATION NUMBER: US/10/738,809
CURRENT FILING DATE: 2003-12-17
PRIOR APPLICATION NUMBER: US/10/344,779
PRIOR APPLICATION NUMBER: PCT/US01/26161
PRIOR PILING DATE: 2003-02-19
PRIOR FILING DATE: 2001-08-21
PRIOR APPLICATION NUMBER: US 60/226,524
PRIOR APPLICATION NUMBER: US 60/230,639
DEFORE FILING DATE: 2000-08-21
PRIOR APPLICATION NUMBER: US 60/230,639
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INFORMATION FOR SEQ ID NO: 143:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 07/472,947
FILING DATE: January 31, 1990
ATTORNEY/AGENT INFORMATION:
NAME: Pabst, Patrea L.
REGISTRATION NUMBER: 31,284
REGISTRATION NUMBER: GMRF114
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: peptide SEQUENCE DESCRIPTION: SEQ ID NO: 143:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (404)-817-8473
TELEFAX: (404)-817-8588
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                   Score 13; DB 16;
Pred. No. 1.4e+06;
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Pred. No. 1.4e+06;
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                                          Length 5;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 10
US-10-394-471B-11
                                                                                                                                                                                                                                                                                                                                                                         US-09-990-762-36
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CURRENT APPLICATION NUMBER: US/10/394,471B

CURRENT FILING DATE: 2003-03-21

PRIOR APPLICATION NUMBER: 60/367,903

PRIOR FILING DATE: 2002-03-26

NUMBER OF SEQ ID NOS: 17

SOFTWARE: Patentin Ver 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local
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Patent No. US20020119498A1
                 SOFTWARE: Pa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Giles-Komar, Jill; Carton, Jill; Scallon, Be TITLE OF INVENTION: ANTI-TNF ANTIBODIES, COMPOSITIONS, FILE REFERENCE: CENO202
CURRENT APPLICATION NUMBER: US/10/394,471B
CURRENT FILING DATE: 2003-03-21
PRIOR APPLICATION NUMBER: 60/367,903
PRIOR FILING DATE: 2002-03-26
NUMBER OF SEQ ID NOS: 17
SOFTWARE: Patentin Ver 3.1
                                     APPLICANT: JOUNG, J. KEITH
APPLICANT: MILLER, JEFFREY
APPLICANT: MILLER, JEFFREY
APPLICANT: PABO, CARL O.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INTERACTION TRAP ASSAYS
FILE REFERENCE: MTV-030.02 (20021-3002)
CURRENT FILING DATE: 2001-11-14
PRIOR APPLICATION NUMBER: U5/95/990,762
CURRENT FILING DATE: 2001-15-16
PRIOR FILING DATE: 2001-05-16
PRIOR FILING DATE: 2000-05-16
                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 5
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              B6.7%; Local Similarity 66.7%; hes 2; Conservation
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Pred. No.
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Pred. No. 1.4e+06;
0; Mismatches 1
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Query Match
Best Local Similarity
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CURRENT ELLING DATE: 2001-11-14
PRIOR APPLICATION NUMBER: US/09/990,762
CURRENT FILING DATE: 2001-05-16
PRIOR APPLICATION NUMBER: 09/858,852
PRIOR FILING DATE: 2001-05-16
PRIOR FILING DATE: 2000-05-16
NUMBER OF SEC ID NOS: 97
SOFTWARE: Patentin Ver: 2.1
SEQ ID NO 40
LENGTH: 6
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Matches
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                                                      PRIOR FILING DATE: 2000-05-16
NUMBER OF SEQ ID NOS: 97
SOFTWARE: PATENTIN VEY: 2.1
SEQ ID NO 41
                                                                                                                                                                                                                                                                                                                                                                  Sequence 41, Application US/09990762 Patent No. US20020119498A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                   APPLICANT: JOUNG, J. KEITH
APPLICANT: MILLER, JEFFREY
APPLICANT: MILLER, JEFFREY
APPLICANT: MILLER, JEFFREY
APPLICANT: PABO, CARL O.
APPLICANT: PABO, CARL O.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INTERACTION TRAP ASSAYS
FILE REFERENCE: MTV-030.02 (20021-3002)
CURRENT PERIOR NOTICE: 100/9990,762
CURRENT FILING DATE: 2001-11-14
ERIOR APPLICATION NUMBER: 09/858,852
PRIOR APPLICATION NUMBER: 09/858,852
PRIOR APPLICATION NUMBER: 60/204,509
PRIOR APPLICATION NUMBER: 60/204,509
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: JOUNG, J. KEIT APPLICANT: MILLER, JEFFR APPLICANT: PABO, CARL O.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Description of Artificial Sequence: Zinc finger OTHER INFORMATION: recognition sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Description of Artificial Sequence: Zinc finger OTHER INFORMATION: recognition sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE
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ORGANISM: Artificial Sequence
                                   ENGTH: 6
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3R, JEFFREY
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66.7%;
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Pred. No. 1.4e
0; Mismatches
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Pred. No. 1.4e+06;
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RESULT 15
US-09-858-852A-36
VS-09-858-852A-36
Sequence 36, Application US/09858852A
Publication No. US20030044787A1
Publication No. US20030044787A1
GENERAL INFORMATION:
APPLICANT: MILLER, JEFFREY
APPLICANT: MILLER, JEFREY
APPLICANT: MILLER, JEFREY
APPLICANT: MILLER, JEFREY
COMPOSITIONS FOR INTERACTION TRAP ASSAYS
FILE REFERENCE: MTV-030.01 (20021-3001)
CURRENT APPLICATION NUMBER: US/09/859,852A
FILING DATE: 2001-05-16
PRIOR APPLICATION NUMBER: 60/204,509
PRIOR FILING DATE: 2000-05-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 6, Application US/09518737

Publication No. US20030008321A1

GENERAL INFORMATION:

APPLICANT: FUKUI, YASUHISA
APPLICANT: NAGATA, SATOSHI
APPLICANT: SHIRAI, RYUICHI
APPLICANT: PHOSPHATIDYLINGSITOL-3,4-DIPHOSPHATE
FILE REFERENCE: 1965/49618
CURRENT APPLICATION NUMBER: US/09/518,737
CURRENT FILING DATE: 2000-03-03
PRIOR APPLICATION NUMBER: JP 1999-250209
PRIOR FILING DATE: 1999-09-03

PRIOR FILING DATE: 1999-09-03

NUMBER OF SEQ ID NOS: 10
SOFTWARE: PATENTIN Ver. 2.1

LENGTH: 6
TYES: PRT
ORGANISM: Mus musculus
US-09-518-737-6
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; OTHER INFORMATION: Description of Artificial Sequence: Zinc finger
; OTHER INFORMATION: recognition sequence
US-09-990-762-41
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Best Local Similarity
Matches 2; Conserv
LENGTH: 6
TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Zinc finger
OTHER INFORMATION: recognition sequence
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Best Local Similarity 66.7
Matches 2; Conservative
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tive 0; Mismatches 1;
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Search completed: December 13, 2004, 23:02:39 Job time: 68.6 secs

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T-cell receptor al
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tubulin alpha chai
hemagglutinin - In
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alkaline serine pr	alphaA-conotoxin P	guanidino kinase M	cysteine proteinas	fibrinogen gamma c	T-cell receptor be	שי	alpha 1-proteinase	Ig heavy chain V r	H+-transporting tw	tyrocidine synthas	Ig heavy chain V r	T-cell receptor J-	T-cell receptor J-	Ig heavy chain CDR	pyrrhocoricin - Py

ALIGNMENTS

Ig heavy chain V region - mouse C;Species: Mus musculus (house mouse) C;Date: 06-Jan-1994 #sequence_revision 01-Dec-1995 #text_change 05-Nov-1999 C;Accession: S36850

A;Status: preliminary
A;Molecule type: nucleic acid
A;Residues: 1-9 <JAC>

A;Cross-references: EMBL:X6'C;Keywords: heterotetramer;

EMBL:X67387; NID:g50113; PIDN:CAA47799.1; PID:e51594; PID:g1333871 etramer; immunoglobulin

Length 9;

Similarity

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T-cell receptor gamma chain (2t.23) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_char
C;Accession: G41946
R;Whetsell, M.; Mosley, R.L.; Whetsell, L.; Schaefer, F.V.; N
MO1. Cell. Biol. 11, 5902-5909, 1991
A;Title: Rearrangement and junctional-site sequence analyses
A;Reference number: A41946; MUID:92049316; PMID:1558619
A;Accession: G41946
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C;Keywords: T-cell
                                                                                                                                             A;Status: preliminary;
A;Molecule type: DNA
A;Residues: 1-9 <WHE>
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                                                                                                                              receptor
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                                                                                                                                                                                                                                                       R.L.; Whetsell, L.; Schaefer, F.V.; Miller, K.S.; Klein,
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Pred. No. 2.8e+05;
0; Mismatches 1;
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RESULT 3

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A; Residues: 1-11 < YAM>
A; Residues: 1-11 < YAM>
A; Experimental source: B lymphocyte
A; Experimental source: Inmunoglobulin
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C;Keywords: T-cell receptor
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A;Title: T cell receptor V gene usage of islet beta cell-reactive A;Reference number: PT0209; MUID:91217621; PMID:1902501
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                                                                                                                                                                                                                                                     A;Reference number: A33098
A;Accession: A33099
                                                                                                                                                                                                                                                                                                                                                                   163K exoantigen -
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; Yamada, M.; Wasserman, R.; Richard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.; Exp. Med. 173, 395-407, 1991
Exp. Med. 173, 395-407, 1991
; Title: Preferential utilization of specific immunoglobulin heavy chain diversity; Reference number: PT0272; MUID:91108337; PMID:1899102
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Matches 2
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                                                                                                                                                                                                     ;Molecule type: protein
;Residues: 1-12 <NIC>
                                                                                                                                                                                                                                                                                                                                                   53K exoantigen - malaria parasite (Plasmodium falciparum)
:Species: Plasmodium falciparum
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;Residues: 1-11 <NAK>
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0; Mismatches 1
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Pred. No. 8.1e+02;
0; Mismatches 1
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Pred. No. 8.
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R.Lehner, P.J...
R.Lehner, P.J...
submitted to the EMBL Data Library, August 1994
submitted to the EMBL Data Library, August 1994
submitted recognition
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PH1611
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submitted to the EMBL Data Library, August
A;Description: Human HLA-A0201 restricted 1
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C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change C;Accession: $47391; $47386
                                                                                                                                                                         A;Title: Molecular characterization of transgene-induced A;Reference number: PH1580; MUID:93301609; PMID:8315387 A;Accession: PH1611
                                                                                                                                                                                                                                                             Ig H chain V-D-J region (wild-type clone 341) - mouse (fragment) C;Species: Mus musculus (house mouse) C;Date: 02-Jun-1994 #sequence_revision 02-Jun-1994 #text_change C;Accession: PH1611
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Cross-references: EMBL:235715; NID:g527525; PIDN:CAA84784.1; C;Keywords: T-cell receptor
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A; Accession: S47391
                                                                                                                                       A; Molecule type: DNA
A; Residues: 1-12 <LEV>
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A; Residues: 1-12 < LEH >
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A; Accession: $47395
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C;Keywords: T-cell receptor
                                                                                                     C; Keywords: immunoglobulin
                                                                                                                        A; Experimental source:
                                                                                                                                                                                                                             R;Levinson, D.A.; Campos-Torres, J.; Leder, J. Exp. Med. 178, 317-329, 1993
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A;Residues: 1-12 <LEH>
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Pred. No. 8.8e+02;
0; Mismatches 1
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Pred. No. 8.8e+02;
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Pred. No. 8.8e+02;
0; Mismatches 1
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R; Lehner, P.J.
submitted to the EMBL Data Library,
A; Description: Human HLA-A0201 rest;
A; Reference number: $47355
A; Accession: $47368
                                      A; Molecule type: mRNA
A; Residues: 1-13 < LEH >
                                                                               A; Status: preliminary
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J. Exp. Med. 174, 1371-1383, 1991
A;Title: T cell receptor genes in a series of class I major histocompatibility complexablelic exclusion and antigen-specific repertoire.
A;Reference number: PH0746; MUID:92078846; PMID:1836010
A;Accession: PH0802
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PH0790
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A; Experimental source: T lympho
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A; Residues: 1-12 < CAS>
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PH0802
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           Cross-references: EMBL: 235693; NID: 9527477; PIDN: CAA84762.1; PID: 9527478
                                                                                                                                                                                                                        -cell antigen receptor VJ junction beta chain - human

*Species: Homo sapiens (man)

*Pate: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 05-Nov-1999
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;Residues: 1-12 <CAS>
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Pred. No. 8.8e+02;
0; Mismatches 1
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                                                                    unidentified 6.3/40K protein [imported] -
[;Species: Oryza sativa (rice)
[;Date: 20-Apr-2000 #sequence_revision 20-
[;Accession: P00700
         R;Komatsu, S.; Kajiwara, H.; Hirano, H.
Theor. Appl. Genet. 86, 935-942, 1993
A;Title: A rice protein library; a data-file
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A;Cross-references: EMBL:X60893
A;Experimental source: T lympho
C;Keywords: T-cell receptor
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A; Reference number: PH0746; MUID:92078846; PMID:1836010
                                                                                                                                                                                                                                                                                                                               T-cell receptor alpha chain (E22 V-alpha-8.F3.4) - mouse (fragment) C;Species: Mus musculus (house mouse) C;Decies: Mus musculus (house mouse) C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997 C;Accession: PH0788
                                                                                                                                                                                      A; Molecule type: mRNA
A; Residues: 1-13 < CAS>
                                                                                                                                                                                                                                                                                          R;Casanova, J.L.; Romero, P.; Widmann, C.; Kourilsky, P.; Maryanski, J.L. J. Exp. Med. 174, 1371-1383, 1991
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C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995
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A; Residues: 1-13 < LEH >
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A; Accession: S47383
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                                                                  Conservative
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20-Apr-2000 #text_change

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separated

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two-dimension

rice (fragment)

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OCC1 enhancer binding protein 70K chain MUCIEBP-70 - human (fragment)
C;Species: Homo sapiens (man)
C;Accession: A59018
R;Abe, M, Smith, C.J.; Larson, C.J.
submitted to the Protein Sequence Database, May 1998
A;Description: Involvement of "Ku-like" proteins in the transcription of MUC1/DF3, a bre A;Reference number: A59018
A;Accession: A59018
A;Accession: A59018
A;Accession: A59018
A;Accession: A59018
A;Molecule type: protein
A;Residues: 1-14 <ABEs
A;Experimental source: breast cancer cell line MCF-7
A;Note: 3-Val was also found
C;Keywords: DNA binding; heterodimer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 15
A59018
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Status: preliminary
A;Molecule type: protein
A;Residuse: 1-13 -KOM5
A;Cross-references: UNIPROT:Q7M1U1
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Search completed: December 13, 2004, 22:32:59 Job time : 16.2 secs
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                                                                                                                                                                                   Query Match 86.7%; Score 13; DB 2; Length 14; Best Local Similarity 66.7%; Pred. No. 1e+03; Matches 2; Conservative 0; Mismatches 1; Indels
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Aac39714 homo sapi
Aab92363 homo sapi
O9gjw1 lagenorhync
O9gjw2 lagenorhync
O9gjw3 lagenorhync
O6ld78 clostridium
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homo sapien
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UF06_MOUSE
ID UF06 MOUS.
C 93644;
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01-CCT-1994 7-UUL-2/
RESULT
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Best Local Similarity
Matches 2; Conserv
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Q9QWT0;
Q9QWT0;
01-MAY-2000 (TrEMBLrel. 13, C
01-MAY-2000 (TrEMBLrel. 13, L
01-DEC-2001 (TrEMBLrel. 19, L
Proteinase 3 (Myeloblastin) (
                                                                                                                                                                                                                Mus musculus (Mouse).
Eukaryota; Metazoa; C
Mammalia; Eutheria; R
NCBI_TaxID=10090;
SEQUENCE FROM N.A.
STRAIN-SV129 D3;
Lutz P.G., Moog-Lutz C., Houzel-Charavel A., C
Submitted (UUN-1998) to the EMBL/GenBank/DDBJ
EMBL; AJ007030; CAA07429.1; -
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MEDILINE=95009907; PubMed=7523108;

Merrick B.A., Patterson R.M., Wichter L.L., He C., Selki "Separation and sequencing of familiar and novel murine preparative two-dimensional gel electrophoresis.";

Electrophoresis 15:735-745(1994).

Electrophoresis 15:735-745(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein is: 5.2, its MW is: 50 KDa.
Direct protein sequencing.
NON TER 8
SEQUENCE 8 AA; 817 MW; A35DD878676B0
                                                                                                                                                                                                                                                                                                                                                                                                           Name=Prtn3;
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus

NCBI_TaxID=10090;
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01-0CT-1994 (Rel. 30, Last sequence update)
05-UTL-2004 (Rel. 44, Last annotation update)
Unknown protein from 2D-PAGE of fibroblasts (
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Rodentia;
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Q9R4F9
Q9R501
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Q9N6I3
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Q25281
Q9GNL7
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Pred. No. 1.8e+06;
0; Mismatches 1
                                                                                                                                                                                                                                                                           Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus.
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                                          Cayre Y.E.;
J databases.
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urine proteins
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Q9r469 aeromonas.
Q9r501 bacillus su
Q9r501 bacillus su
Q9prx1 struthio ca
Aab20893 clostridi
Q9n613 toxoplasma
Q9gn16 drosophila
Q9r4f8 aeromonas.
Q9f4f8 yersinia pe
P37362 yersinia pe
P37362 yersinia pe
P37362 pyrrhocoris
Q25281 leishmania
Q9gn17 drosophila
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Result No.

Score

Length

ВB

Description

score greater than or equal

UniProt_02:*
1: uniprot_sprot:*
2: uniprot_trembl:*

No. is the number of results predicted by chance to have a greater than or equal to the score of the result being printed,

derived by analysis of the total score distribution

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Q9SBIA Q47893 Q47893 Q47893 PH2 PRUSE Q7LDD2 Q6LDD2 Q6LD19 Q6LC19 Q62645 Q911VQ4 Q912H6 Q912H6 Q912H1 Q912B1 QAB92363 Q9GJW1 Q9GJW1 Q9GJW1 Q9GJW3 Q9GJW3

Q61c19 062645 Q71vq4

Q71gd2 Q6jdk5 Q9smv3

Q47892 Q47893 P29264

P80620 Q98814

ADFB TENMO P82432 Q7M1U1 Q7YOR1 PH3 PRUSE UC14 MAIZE

Q7mlul

AAO63472 Q9UC46 Q7YP62 UF06_MOUSE

Q8ney9 homo sa Aao63472 homo Q9uc46 homo sa

sapien mo sapi

Minimum Maximum

BB

seq

length: length:

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number of

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parameters:

1825181 segs, 575374646 residues

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 sp

summaries

Title: Perfect

score:

US-10-068-905-6 15 GXYX 4

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protein

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protein search, using

æ model GenCore version 5.1.6 Copyright (c) 1993 - 2004 Compugen Ltd

December 13, 2004, 22:00:41; Search time 83.8 Seconds (without alignments) 27.464 Million cell updates/sec

Scoring table:

BLOSUM62 Gapop 10.0 ,

Gapext

66.7%;

Pred. No.

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RESULT 3
GREEP9
ID ORNE
ORNE
DT 01-0
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DE TYPE
GN Name
OC Huka
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                                                               SOURCE SERVING SERVING
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01-OCT-2002
01-OCT-2002
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Submitted (DEC-2002) to the EMBL/GenBank/DDB:
EMBL; AY037552; AAK86868 1; -.
EMBL; AY037953; AAO65472.1; -.
GO; GO:0005882; C:intermediate filament; IEA.
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01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
05-UUL-2004 (TrEMBLrel. 27, Last annotation update)
Type II hair-specific keratin (Type II hair keratin)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
Bairwa N.K., Bamezai
Submitted (MAY-2001)
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Mammalia; Eutheria;
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                                                                                            NON_TER
                                                                                                                                                     "Study of promoter polymorphism of type II hair keratin gene hHb6."; "Study of promoter polymorphism of type II hair keratin gene hHb6."; Submitted (DBC-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; AY203963; AA063472.1; -.
                                                                                                                                                                                                                                                                                                                                                              Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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02-MAR-2004 (TrEMBLrel. 27, Last annotation
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01-JUN-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Neutrophil inhibitor peptide, NIP=POLYMORPHONUCLEAR
inhibitor peptide.
                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
Liu Y., He X., Xu L.,
Submitted (JUN-2003) t
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01-OCT-2003 (TrEMBLrel. 25, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation updat
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"Characterization of a neutrophil inhibitor peptide harvested frought and prochial lavage: homology to influenza A nucleoprotein.";
Am. J. Respir. Cell Mol. Biol. 15:207-215(1996).
GO; GO:000576; C:extracellular; NAS.
GO; GO:000526; P:anti-inflammatory response; NAS.
SEQUENCE 11 AA; 1262 MW; 951A1C3279C9DB45 CRC64;
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Mammalia; Eutheria;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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01-JUN-2000 (TrEMBLrel. 24, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
32 kDa cell wall protein (Fragment).
Nicotiana tabacum (Common tobacco).
Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta; Wagnoliophyta; eudicotyledons; core eudicots;
lamids; Solanales; Solanaceae; Nicotiana.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antidiuretic peptide.";
Peptides 24:27-34(2003).

Peptides 24:27-34(2003).

Peptides 24:27-34(2003).

Peptides 24:27-34(2003).

Peptides 24:27-34(2003).

Secretion by Malpighian tubul.

CGMP as second messenger. May function as an antidiuretic composition.

Secreted.

TISSUE SPECIFICITY: Brain. Appears to be mainly expressed pairs of bilaterally symmetrical cells in the protocerebru pairs of bilaterally symmetrical cells in the protocerebru MASS SPECIFIOMETRY: MW=1560.33; METHOD=MALDI; RANGE=1-13; NOTE-REF1.

NOTE-REF1.

Direct protein sequencing; Hormone; Neuropeptide.

SEQUENCE 13 AA; 1562 MW; 0240A4504B8A632B CRC64;
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NCBI_TaxID=7067;
                                                                                                                                                                                                                Cell wal
                                                                                                                                                                                                                                                                   STRAIN-cv. PETIT HAVANA;
Blee K.A., Bonham V.A., Mitchell G.P., I
Wojtaszek P., Bolwell G.P.;
"Protecomic study of secondary cell wall
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eigenheer R.A., Wiehart U.M., N
Hull J.J., Schooley D.A.;
"Isolation, identification and
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P83109;
10-OCT-2003 (Rel. 42, Created)
10-OCT-2003 (Rel. 42, Last seq
05-JUL-2004 (Rel. 44, Last ann
                                                                                                                                                                                    SEQUENCE
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                                                                  Conservative
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iehart U.M., Nicolson
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1307 MW;
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Pred. No. 8.3e+03;
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RESULT 8 P82432

P82432

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Q7YOR1,
01-OCT-2003 (TrEMBLrel. 25, Create
01-OCT-2003 (TrEMBLrel. 25, Last s
01-MAR-2004 (TrEMBLrel. 26, Last a
Gamma-glutamylcysteine synthetase
                                                                                                                          SEQUENCE FROM N.A.

**Machter A., Steininger H., Rausch T., Bogs in Wachter A. (MAY-2003) to the EMBL/GenBank/DDI EMBL; AJ564376; CAD91995.1; -

GO; GO:0004357; F:glutamate-cysteine ligase GO; GO:0016874; F:ligase activity; IEA.
                                                                                                                                                                                                                                                                        Brassica juncea (Leaf mustard) (Indian mustard). Embryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Amgnoliophyta; eudicotyledons; core eudicots; rosid: eurosids_II;_Brassicales; Brassicaceae; Brassica.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Komatsu S., Kajiwara H., Hirano H.;
"A rice protein library; a data-file of two-dimensional electrophoresis.";
Theor. Appl. Genet. 86:935-942(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unidentified 6.3/40K protein (Fragment).
Oryza sativa (Rice).
Eukaryota, Viridiplantae, Streptophyta, Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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(TrEMBLrel. 26, Last sequence up)
(TrEMBLrel. 26, Last annotation)
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                 86.7%;
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66.7%;
                                                                        MW;
                                                                                                                                                                                 Rausch T., Bogs J.;
EMBL/GenBank/DDBJ
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Last annotation update)
hetase (EC 6.3.2.2) (Fragment).
0;
Score 13; DB Pred. No. 8.3e 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                          Created)
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Pred. No.
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                                                                        CRC64;
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                                   Length 13;
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                                                                                                                                                                                                                                                                                              eudicots; rosids;
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Best Local
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 Best Local Similarity
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P80620;
01-OCT-1996
01-OCT-1996
05-JUL-2004
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01-DEC-1992
01-DEC-1992
05-JUL-2004
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01_DEC-1992 (Rel. 24, Last sequence update)
05_DUL-2004 (Rel. 44, Last annotation update)
05_DUL-2004 (Rel. 44, Last annotation update)
Prunasin beta-glucosidase IIB (EC 3.2.1.118) (Prunasii
isozyme IIB) (PH IIB) (Fragment).
Prunus serotina (Black Cherry).
Eukaryota, Viridiplantae, Streptophyta, Embryophyta; Core eugermatophyta, Magnoliophyta; eudicotyledons; core eugersids I, Rosales, Rosaceae, Amygdaloideae, Prunus.
                                                                                                                                                                                                                                                                                                                                            MAIZE
                                                                                    Li C.P., Swain E., Poulton J.E.;
"Prunus serotina amygdalin hydrolase and prunasin hydrolase.";
Plant Physiol. 100:282-290(1992).
-i- CATALYTIC ACTIVITY: (R)-prunasin + H(2)O = mandelonitrile +
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE=Seed;
Li C.P., Swa
                                           NON TER
                                                                                                                                        Touzet P., Riccardi F., Morin (Pernollet J.-C., Zivy M., de Vi
"The maize two dimensional gel
                                                                                                                                                                                                 ___uc; Pa
NCBI_TaxID=4577;
[1]
                                                                                                                                                                                                                    Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD clade; Panicoideae; Andropogoneae; Zea.
                                                                                                                                                                                                                                                                          01-OCT-1996 (Rel. 34, Created)
01-OCT-1996 (Rel. 34, Last sequence update)
05-UTL-2004 (Rel. 44, Last annotation update)
Unknown protein from 2D-PAGE of etiolated coleoptile
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                                SEQUENCE
                                                               Direct protein sequencing.
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SUBUNIT: Monomer.
DEVELOPMENTAL STAGE: Absent fro
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86.7%;
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                                  CRC64;
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udicots; rosids;
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Q47892;
01-NOV-1996
01-NOV-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=86233345; PubMed=3086870; Conley P.B., Lemaux P.G., Lomax T.L., Grossman A.R.; "Genes encoding major light-harvesting polypeptides are c. "genes encoding the cyanobacterium Fremyella diplosiphon."; Proc. Natl. Acad. Sci. U.S.A. 83:3924-3928(1986).
EMBL; M13216; AAA24871.1; -.
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Biochem. J. 302:15-21(1994).
GO; GO:0008422; Fiseta-glucosidase activity; I
SEQUENCE 15 AA; 1667 MW; 46C40370DC8C730B
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-UN-2003 (TrEMBLrel. 24, Last annotation update)
Beta-D-glucoside glucohydrolase (SC 3.2.1.21) (Fragment).
   Q47893
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Fremyella diplosiphon (Calothrix PCC 7601).
Bacteria; Cyanobacteria; Nostocales; Microchaetaceae; Fremyella.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bukaryota; Viridiplantae; Streptophyta; En
Spermatophyta; Magnoliophyta; Liliopsida;
PACCAD clade; Panicoideae; Andropogoneae;
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Zea mays (Maize)
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Search completed: December 13, 2004, 22:31:36 Job time: 83.8 secs
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                                                                                                                                                                                                                                     SEQUENCE FROM N.A.

MEDLINE=8623345; PubMed=3086870;

CONLEY P.B., Lemaux P.G., Lomax T.L., Grossman A.R.;

"Conley P.B., Lemaux P.G., Lomax T.L., Grossman A.R.;

"Genes encoding major light-harvesting polypeptides are clustered on the genome of the cyanobacterium Fremyella diplosiphon.";

Proc. Natl. Acad. Sci. U.S.A. 83:3924-3928(1986).

EMBL; M13217; AAA24880.1; -.

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SEQUENCE 15 AA; 1528 MW; 7FF2F65518F493D4 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                           Q47893;
01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-1998 (TrEMBLrel. 08, Last annotation update)
Beta-phycocyanin (Fragment).
Fremyella diplosiphon (Calothrix PCC 7601).
Bacteria; Cyanobacteria; Nostocales; Microchaetaceae; Fremyella.
NCBI TaxID=1197;
[1]
                                                                                                                   1 GXY 3
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17.936 Million cell updates/sec
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Copyright (c) 1993 - 2004 Compugen Ltd.
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Abp59244	Abj18639	Abb75621	Aae16333	Abp67642	Abp67641	Aab66249	Aab66243	Aab08397	Aay68375	Aay68369	Aay44679	Aay43367	Aay43373	Aaw95236	Aaw95230	Aay41826	Aay41820	Aar72699	Aar/2549
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ALIGNMENTS

RESULT 1 AAP61656

25-MAR-2003 03-OCT-2002 21-AUG-1991

(revised)
(revised)
(first entry)

AAP61656;

AAP61656 standard; protein; 4 AA.

Sequence of peptide which inhibits cyclic-nucleotide independent protein kinase activity and mammalian cell growth.

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Claim 1; Col 4; 9pp; English.
                                                     Inhibition of tumour cell growth - using peptide and aminoacid halo:methyl\ ketone(s).
                                                                         WPI; 1986-118872/18.
                                                                                     Kettner CA, Racker E;
                                                                                                              16-NOV-1983;
                                                                                                                          16-NOV-1983;
                                                                                                                                       15-APR-1986.
                                                                                                                                                    US4582821-A.
                                                                                                                                                                      Misc-difference
                                                                                                                                                                                         Key
                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                               Cell growth inhibitor; tumour cell growth inhibitor.
                                                                                                                                                                                   Misc-difference
                                                                                                  (DUPO ) DU PONT DE NEMOURS & CO
                                                                                                               83US-00552255.
                                                                                                                          83US-00552255.
                                                                                                                                                               /label= Tyr-CH2Cl
                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                           label= Carbobenzoxy-Phe
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The cpds. of the invention inhibit protein phosphorylation. The inventor claim a process for inhibiting the growth of tumour cells in a medium which comprises contacting the cells with a cpd. of formula (AAP61654-P61661) or a physiologically acceptable salt. (Updated on 03-OCT-2002 the add missing OS field.) (Updated on 25-MAR-2003 to correct PA field.)

The inventors

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RESULT 2
AAP71
AAP7
XX APP
AAP7
XX APP
DT 15-M
DT 15-M
DT 15-M
XX Picc
XX Synt
XX Synt
XX Picc
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AAR55333
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Matches 2; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                      This peptide is useful as part of a peptide/halo-methyl ketone cpd., for treating picornavirus, egpolio- or rhinovirus infections. It inhibits the processing of picornavirus capsid proteins by virus encoded proteases. See AAP71301-12. See also US4652552. (Updated on 25-MAR-2003 to correct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAP71313 standard; protein; 4 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Treating picorna-virus infection with peptide halo: methyl ketone cpds. esp. for treating polio virus and rhino virus infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-JAN-1987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Picornaviridae; poliovirus; rhinovirus; antiviral agent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peptide component of cpd. for treating picornavirus infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-MAR-2003
15-MAY-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAP71313;
                                                                                                                                                                                                                                                                                                                                         Sequence 4 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 4; 10pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1987-036897/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kettner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-AUG-1984;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-AUG-1984;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US4636492-A
                    Inhibitor of tyrosine kinase.
                                                 25-MAR-2003
30-JAN-1995
                                                                                                 AAR55333;
                                                                                                                                AAR55333 standard;
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                                                                                                                                peptide; 4 AA
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1.7e+06;
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27-OCT-1993;
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Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1994-183419/22.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-NOV-1993;
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13-NOV-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 4 AA;
                                                                                                                                                                                                                                                                                                                                  AAR75499;
                                                        06-DEC-1993;
                                                                                                                      15-JUN-1995.
                                                                                                                                                     WO9516033-A1
                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                 AAR75499 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (WARN ) WARNER LAMBERT
                        (CIBA ) CIBA
                                                                                     16-NOV-1994;
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2; Conserv
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                           GEIGY AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mcnamara
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The sequence is that of a protein tyrosine kinase inhibitor contg. D- Tyr or tetrafluoro-Tyr residues. The peptide can be used to inhibit TK-mediated signal transduction and is useful for controlling proliferative diseases, e.g. cancer, psoriasis and to treat viral, inflammatory, allergic and cardiovascular diseases. See also AAR55302-39. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New peptide inhibitors of protestrafluoro-Tyr residue, for infection, inflammation etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PTK; signal transduction; proliferation; cancer; psoriasis; viral;
inflammatory; allergic; cardiovascular; diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 7; Page 36; 44pp; English.
                                                                                                                                                                                                                                           C5a receptor-antagonist core
                                                                                                                                                                                                        C5a receptor-antagonist;
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(first entry)
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93US-00139913.
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93US-00162591
                                         94WO-IB000359
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "Tyr(2,3,5,6-F4) (D/L)"
                                                                                                                                                                                                                                                                                                                                                                                             peptide; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     86.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 13; DB
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                                                                                                                                                                                                                                                tetrapeptide.
                                                                                                                                                                                                        antiinflammatory.
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The present invention describes a human leukocyte antigen (HLA) (e.g. soluble HLA) ligand database assembled by a process which involves providing a computer system capable of storing HLA data as a database, producing HLA having ligands loaded on it, isolating the loaded ligands from HLA, sequencing the loaded ligands to obtain HLA ligand data, and populating the database with HLA ligand data. Also described: (1) accessing HLA (e.g. sHLA) data stored in a database, and (2) a computer system for HLA (e.g. sHLA) ligand database. The database is populated with HLA ligand sequences, motifs, extended motifs, submotifs, ligands.
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10-OCT-2001; :
18-DEC-2001; :
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displays C5a
field.)
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                                                                                                                                                                                                                                                             (HILD/)
                                                                                                                                                                                                                                                                                                                                              21-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                              HLA ligand;
MHC ligand;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Polypeptide receptor antagonists to C5a contain an essential tetrapeptide (AAR75498 or AAR75499) or core tripeptide (DGA) displays C5a blocking activity. (Updated on 25-MAR-2003 to co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C5a receptor antagonists having compsns. to treat C5a-mediated
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                                                                                                                                           leukocyte antigen ligand database assembled by producing HLA having d ligands, isolating and sequencing loaded ligands to obtain HLA data and populating database in computer system with ligand data.
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                                                                                                                                                                                                                                                                                                                                                                                                                                           human leukocyte antigen; major histocompatibility
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sHLA molecule ligand related peptide
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Pred. No. 1.7e+06;
0; Mismatches 1;
                                                                                                                                                                                                                         Hickman
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                                                                                                                                                                                                                                                                                                                                                                                                                                           predictive algorithm; database;
complex; viral; bacterial; tumo
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RESULT 6
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10-OCT-2001;
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(PRIL/)
(HICK/)
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ligand; major
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2; Conserv
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Pred. No. 1.7e
0; Mismatches
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complex; viral; bacterial; tumo;
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1.7e+06;
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The present invention describes a human leukocyte antigen (HLA) (e.g. soluble HLA) ligand database assembled by a process which involves providing a computer system capable of storing HLA data as a database, producing HLA having ligands loaded on it, isolating the loaded ligands from HLA, sequencing the loaded ligands to obtain HLA ligand data, and populating the database with HLA ligand data. Also described: (1) accessing HLA (e.g. sHLA) data stored in a database; and (2) a computer system for HLA (e.g. sHLA) ligand database. The database is populated with HLA ligand sequences, motifs, extended motifs, submotifs, ligands

Human leukocyte antigen ligand database assembled by producing HLA loaded ligands, isolating and sequencing loaded ligands to obtain I ligand data and populating database in computer system with ligand

HLA having ain HLA

data.

WPI; 2002-732755/79

Disclosure; Fig 5; 118pp; English.

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RESULT 7
ARG96817
ARG96817
ARG9682
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ARG98
AXX ARG98
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The invention describes a method of producing soluble human leukocyte antigen (HLA) molecules (SHLA) in cell pharm involving amplifying HLA allelic DNA by pCR using a locus specific primer to produce truncated a pCR product (PI), inserting P1 into mammalian expression vector; electroporating the plasmid into a host cell; inoculating the cell pharm with the host cell such that cell pharm produces sHLA. A multimeric HLA complex (I) is useful for testing functionality of peptide ligands bound
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-DEC-2000; 2000US-0256409P.
18-DEC-2000; 2000US-0256410P.
10-CCT-2001; 2001US-00974366.
                                                                                                                                                                                                                                                                                                                                Producing soluble human leukocyte antigen (HLA) in cell pharm useful studies of peptide loading for characterizing human immune responses involves using HLA allelic cDNA or genomic DNA as starting material.
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; Pred. No. 1.7e
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1.7e+06;
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development; MHC;
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cc by at least two soluble HLA molecules. (I) can be tested for its ability to serve as ligands for cytotoxic T lymphocytes (CTLs) and induce immune cresponses in humans. (I) is useful for studying T cell responses to pathological conditions such as viral infections and cancer, and for condulating the human immune system to induce tolerance in autoimmune conditions and responses to molecules produced are useful for studies of peptide loading (i.e., in concerned MHC molecules, naturally loaded peptide loading (i.e., in concerned MHC molecules, naturally loaded peptides can be eluted from the consensuant of structural and functional impact of HLA class I colymorphism. The molecules are also useful to generate ligands and hence conjumorphism. The molecules are also useful to generate ligands and hence conjumorphism. The molecules pools extracted from series of distinct yet conjumorphism. The molecules of series of HLA-B15 allotypes; compare the different ligand maps to cidentify potentially shared elements; and characterise the elements conjumorphism insect/bacterial cells such that milligram or greater constitutes of an individual class I molecule can be obtained. This sequence represents a HLA (human leukocyte antigen) peptide ligand was to characterise of an individual class I molecule can be obtained.
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18-DEC-2000;
10-OCT-2001;
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                                            producing soluble human leukocyte antigen (HLA) in cell pharm useful studies of peptide loading for characterizing human immune responses involves using HLA allelic cDNA or genomic DNA as starting material.
                                                                                                                                                                                        (HILD/)
(PRIL/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABG97024 standard; peptide; 4
                                                                                                                     WPI; 2002-698563/75
                                                                                                                                                                                                                                                                                                                                                                                  WO200262846-A2
                Disclosure;
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                                                                                                                                                                                        HILDEBRAND W H. PRILLIMAN K R.
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                Page
                                                                                                                                                                                                                                         ; 2000US-0256409F.
; 2000US-0256410F.
; 2001US-00974366.
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                                                                                                                                                       Prilliman
                183; 300pp; English.
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66.7%;
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Pred. No.
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1.7e+06;
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RESULT 9
ABG96697
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Soluble human leukocyte antigen; HLA; sHLA; cell pharm; multimeric HLA complex; cytotoxic Tlymphocytes; CTL; immune responsal infection; cancer; autoimmune disease; vaccine development; major histocompatibility complex; diagnostic development; HLA-B15 allotype; ligand.
Hildebrand WH,
                                                                                                                                                                                      18-DEC-2000; 2000US-0256409P.
18-DEC-2000; 2000US-0256410P.
10-OCT-2001; 2001US-00974366.
                                                                          (HILD/)
(PRIL/)
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                                                                                                                                                                                                                                                                                                                                                   18-DEC-2001; 2001WO-US049744.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-DEC-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 4
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                                                                      HILDEBRAND W H.
PRILLIMAN K R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N
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2; Conserv
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    Prilliman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      86.7%;
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Pred. No. 1.7e+06;
0; Mismatches 1;
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development; MHC;
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(SKUB/)

SKUBITZ K M. SKUBITZ A P

28-FEB-2001; 2001US-0272113P

27-FEB-2002; 2002WO-US005720

06-SEP-2002

Homo sapiens.

Human; CD66; CEACAM; cytc antibacterial; virucide; immune system; autoimmune

inflammatory disease; transplantation;

Human

CD66 family modulating peptide SEQ ID

No

CEACAM; cytostatic; antiinflammatory;

immunomodulator;

virucide; gene therapy; vaccine; neutrophil; autoimmune disease; cancer; infection; bacte

10-DEC-2002

(first entry)

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RESULT 10
ABP67652
ID ABP677
XX ABP677
XX ID IO-DE
XX Human
XX Human
XX Human
XX Human
XX Antilb
XX Infla
XX In
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Best Local S
Matches 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 13; DB 5; Le
Pred. No. 1.7e+06;
0; Mismatches 1;
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RESULT 11
ABP67653
ID ABP67
XX ABP67
XX ABP67
XX Huma
DE Huma
XX Infi

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New peptide from a surface exposed region of a CD66 family member, useful for modulating the function of CD66 family members, e.g. activation of neutrophils, for treating or diagnosing autoimmune diseases or cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antibacterial; virucide; gene therapy; vaccine; neutrophil;
immune system; autoimmune disease; cancer; infection; bacterial; virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human CD66 family modulating peptide SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-DEC-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CD66; CEACAM; cytostatic; antiinflammatory; immunomodulator;
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2; Conserv
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SKUBITZ A P N.
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                                                                                                                                                                                         Skubitz
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.7e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 5;
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RESULT 12
AAUB4247
ID AAUB4
XX AAUB4
XX AAUB4
XX Caspa
XX Caspa
XX Caspa
KW apopt
KW apopt
KW wyocas
KW myocas
KW mpi he-W
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Best Local &
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to an isolated peptide (I) from a surface exposed region of a CD66 family member (ABP67130-ABP67990). The peptide modulates: (a) activation of neutrophils; (b) activation or inhibition of T-cells, B-cells, INK cells, isolativitic cells or other immune system cells; (c) proliferation and/or differentiation of the above cells, including epithelial cells; (d) homotypic and/or heterotypic adhesion among CD66 family members; and (e) adhesion of CD66 family members; and (e) adhesion of CD66 family members; and (or their ligands, such as activation of neutrophils and activation, inhibition, proliferation and/or differentiation of the immune cells. The peptides may also be used in treating or diagnosing autoimmune diseases, cancer, infections (e.g. bacterials and for immunication of these and for immune cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and as a vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Caspase; apoptosis; GAP SH3 binding protein; HCD2; AOP-1; cancer; apoptosis-associated disease; hyperproliferative disease; AIDS; degenerative disease; autoimmune disease; Alzheimer's disease; viral infection; acquired immunodeficiency syndrome; vascular disease; myocardial infarction; oxidative stress; ischaemic stroke; arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 4 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            therapies and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Caspase cleavage site #11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAU84247 standard; peptide;
                                                                                                                                Novel apoptosis-associated and/or -modified protein useful as target for the diagnosis, prevention or treatment of apoptosis-associated diseases
                                                                                                                                                                                          WPI; 2002-098088/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               heart failure; Parkinson's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   myocardial infarction; oxidative stress; ischaemic stroke; arthrit
heart failure; Parkinson's disease; amyotrophic lateral sclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAU84247;
                                                                            Claim
                                                                                                              particularly neurodegenerative diseases.
                                                                                                                                                                                                                              Rudel
                                                                                                                                                                                                                                                                                                      16-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                            15-JUN-2001; 2001WO-EP006780
                                                                                                                                                                                                                                                                                                                                                                                                 20-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200196873-A2
                                                                                                                                                                                                                                                                  (PLAC ) MAX PLANCK GES FOERDERUNG WISSENSCHAFTEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity hes 2; Conserv
                                                                                                                                                                                                                              Ţ,
                                                                        30; Page 62; 86pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 GXY 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GSY 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 30; 96pp;
                                                                                                                                                                                                                              Thiede B,
                                                                                                                                                                                                                                                                                                      2000EP-00112813.
2000EP-00125013.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  for immunisation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      86.7%;
                                                                                                                                                                                                                            Machuy N;
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Pred. No. 1.7e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <u>.</u>.
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The invention relates to an apoptosis-associated and/or -modified protein (I) selected from GAP SH3 binding protein, HCD2 and AOP-1 (not defined) or its proteolytic fragments. (I) is useful as target for the diagnosis,

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1 GXY 3

Query Match Best Local S Matches 2

2 Similarity

Conservative

0

86.7%; 66.7%;

Score 13; DB Pred. No. 1.7e 0; Mismatches

DB 6; HC... 1.7e+06; 1;

Length 4; Indels

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Gaps

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RESULT 13
ABJ18638
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Best Local S
Matches 2
              preventing tumours and cancers (e.g. gastric cancer, breast cancer and prostate cancer). The present amino acid sequence represents an erbB receptor peptide antagonist of the invention
                                                                                                                                                                                                                                                                                                  New peptide antagonists against erbB receptors, useful treating tumors or cancers (e.g. lung adenocarcinomas,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-APR-2001;
03-AUG-2001;
                                                                                                                                                                                                                        Claim 1; Page 82; 115pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vaccine; erbB receptor antagonist; tumour; cancer; gastric cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ErbB receptor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Greene MI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-APR-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VINU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard; peptide;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4 AA;
                                                                                                                                                                                                                                                                          cancer) in humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Zhang H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2002WO-US011211.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PENNSYLVANIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2001US-0282037P
2001US-0309864P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       caspase cleavage sites of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     86.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4
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Pred. No. 1.7e+06;
0; Mismatches 1
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4 AA

ADJ81984;

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for preventing or breast carcinomas
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               RESULT 15
ADJ81984
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ABP59243
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                                                                                                                           Query Match
Best Local Similarity
Matches 2; Conserv
                                                                                                                                                                                                        The present invention relates to novel antibodies, which have specificit for activated cell surface receptors (erbB, TMF and immunoglobin gene superfamily: 1959) and bind to an epitope (ABP5923). The antibodies are useful for preventing transformation of a normal cell inta tumour cell in a patient who had a tumour with tumour cells that express activated erbB receptors on the cell surface surgically removed or in whom the tumour has entered remission, for treating a patient suffering from a cancer characterised by tumour cells that have pl85 on their cell surfaces, or for treating a patient suffering from an erbB tumour. The antibodies were produced by tumourising a suitable host with peptide (ABP59223-ABP59259), or its reverse sequence
 ADJ81984
                                                                                                                                                                                  Sequence 4 AA;
                                                                                                                                                                                                                                                                                                                                                                                                          Novel antibody for preventing, treating, imaging or diagnosing tumor, has specificity for activated cell surface receptor and binds specifically to an epitope that does not comprise phosphotyrosine residue.
                                                                                                                                                                                                                                                                                                                                                                                     Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Greene MI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-AUG-2001; 2001US-0309864P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antibody; cytostatic; vaccine; cell simmunoglobin gene superfamily; IgSF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO2003012072-A2
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                                                                                                                                                                                                                                                                                                                                                                                   53; Page 88; 114pp;
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standard; peptide; 4
                                                                                                GXY 3
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                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Zhang H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first
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                                                                                                                                       86.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Richter M,
                                                                                                                                                                                                                                                                                                                                                                                   English.
                                                                                                                                       Score 13; DB 6;
Pred. No. 1.7e+06;
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                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             surface
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Search completed: December 13, 2004, 22:22:40 Job time : 81 secs
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                                                                                                 Query Match
Best Local Similarity 66..
"--ches 2; Conservative
                                                                                                                                                                                                           The invention relates to novel nucleic acid polymerases from the thermophilic organism Thermus brookianus or mutants having a mutation that decreases 5'-3' exonuclease activity or reduces discrimination against dideoxymucleotide triphosphates. The nucleic acid polymerases, isolated nucleic acids encoding the polymerases, vectors, bost cells and kits are useful for DNA synthesis, primer extension, DNA sequencing, reverse transcription, or DNA and RNA amplification procedures. The nucleic acids are also useful for the recombinant expression of the polymerase polymerase polypeptides. The nucleic acid polymerases and nucleic acids encoding DNA polymerase I activity. This sequence corresponds to a peptide of a polymerase of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nucleic acid polymerase; thermophile; mutation; exonuclease activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  New Thermus brockianus nucleic acid polymerases, useful for DNA synthesis, primer extension, DNA sequencing, reverse transcription, DNA and RNA amplification procedures.
                                                                                                                                                                                   Sequence 4 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Thermus brockianus nucleic acid polymerase peptide fragment #2.
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                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; SEQ ID NO 18; 74pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rozzelle J,
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                                                                                          1 GXY 3
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                                                                                                                                         86.7%;
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                                                                                                                          Score 13; DB 7; 1
Pred. No. 1.7e+06;
0; Mismatches 1
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Result
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Maximum DB seq
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score
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length: 2000000000
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*
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               US-07-962-569A-6
US-08-292-968-30
US-08-292-968-30
US-08-618-696-3
US-08-618-696-14
US-08-467-974-36
US-08-467-536-36
US-08-467-536-36
US-08-467-536-31
US-08-467-976-36
US-08-467-976-36
US-08-467-976-36
US-09-032-514-36
US-09-082-514-36
US-09-082-514-36
US-08-014-979-5-14-3
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US-08-014-979-5-14-3
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US-09-301-085-199
PCT-US93-01669-41
PCT-US95-04589-199
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                                                      Sequence 49, Appl
Sequence 2, Appli
Sequence 2, Appli
Sequence 199, App
Sequence 199, App
Sequence 199, App
Sequence 199, App
Sequence 30, Appli
Sequence 30, Appli
Sequence 14, Appl
Sequence 31, Appli
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Sequence 31, Appli
Sequence 36, Appli
Sequence 37, Appli
Sequence 38, Appli
Sequence 31, Appli
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US-08-798-897-49
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28,	8,		Sequence 19, Appl	Patent No. 5200320	ď	Sequence 12, Appl	12,	e 12,	132	•	-	-	Sequence 21, Appl	•	32,	31,	•

ALIGNMENTS

US-08-798-897-49 US-08-798-897-49 Patent No. 5799201 Patent No. 5799201 PATENT NO. 15799201 PATENTION GUASTELLA, John PITLE OF INVENTION: Genes Coding For Bcl-y, a Bcl-2 TITLE OF INVENTION: Homologue UNMER OF SEQUENCES: \$3 CORRESPONDENCE STERNE, KESSLER, GOLDSTEIN & FOX P.L.L.C. STREET: 1100 New York Avenue, N.W., Suite 600 STATE: DC COUNTRY: Washington STATE: DC COUNTRY: USA ZIP: 20005 COMPUTER READABLE FORM: MEDIUM TYPE: Ploppy disk COMPUTER READABLE FORM: MEDIUM TYPE: Ploppy disk COMPUTER: IBM PC compatible OPERATION SYSTEM: PC-DOS/NS-DOS SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION UNMER: US/08/798.897 FILING DATE: PS-NATY 11, 1997 ATTORNEY/ACENT INFORMATION: APPLICATION NUMBER: US/08/798.897 FILING DATE: PS-NATY 11, 1997 ATTORNEY/ACENT INFORMATION: TELEPHONE: 202-371-2540 INFORMATION FOR SEQ ID NO. 49: SEQUENCE CHARACTERISTICS: LENGTH: 4 amino acids TYPE: anino acids TYPE: POSTANDENESS: not relevant TOPOLOGY: linear MOLECULE TYPE: peptide US-08-798-897-49 Query Match Best Local Similarity 66.7%; Pred: No. 3.5e+05; Natches 2; Conservative 0; Mismatches 1; Indels 0; Gaps Cy 1 | AX | 3

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                                                                                                                                                                                                                                                                                      Sequence 2,
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 4 amino acids
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                                                                                                                                                                                                                                                                          tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FO-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Lerner, David, Littenberg, Krumholz & Mentlik STREET: 600 South Avenue West
                                                                                                                                                                                                          APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Foley, Shawn P.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 908-654-5000
                                                                        APPLICANT: van Heeke, Gino
TITLE OF INVENTION: C5a Receptor Antagonists Having
TITLE OF INVENTION: Substantially No. 5837499Agonist Activity
NUMBER OF SEQUENCES: 67
CORRESPONDENCE ADDRESS:
ADDRESSES: Lerner, David, Littenberg, Krumholz & Mentlik
STREET: 600 South Avenue West
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: C5a Receptor Antagonists Having TITLE OF INVENTION: Substantially No. 5807824Agonist Activity NUMBER OF SEQUENCES: 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: peptide
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COMPUTER READABLE FORM:
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nes 2; Conserv
                                                     STREET: 600 Sout
CITY: Westfield
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CLASSIFICATION: 435
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               COUNTRY: U
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o. 5807824
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Galakatos, Nicholas G.
Schmitz, Albert
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Pred. No. 3.8e
0; Mismatches
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US-08-978-523-49
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                                                                                                                                       COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/978,523
FILING DATE: herewith
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/798,897
FILING DATE: February 11, 1997
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Esmond, Robert W.
REGISTRATION NUMBER: 33,893
REGISTRATION NUMBER: 33,893
REGISTRATION NUMBER: 1483.0140002
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Best Local Similarity
Matches 2; Conserv
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GENERAL INFORMATION:
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                                                                      TELEPHONE: 202-371-26
TELEFAX: 202-371-2540
INFORMATION FOR SEQ ID NO:
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NAME: Foley, Shawn P.
TELECOMMUNICATION INFORMATION:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                           TELECOMMUNICATION INFORMATION:
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                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: >- TELEFAX: >- TELEFAX: | >- TELEFAX: | | |
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Washington
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TYPE: amino acid
                                    ENGTH:
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5883229
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4 amino acids
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                                    4 amino acids
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Homologue
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66.7%;
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Pred. No.
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US-08-978-523-49
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Best Local (
                                                                                                                  Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA.
ZIP: 02110-2904
COMPUTER READABLE FORM:
COMPUTER: FLODDY disk
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
COMPUTER: TEM PC DOS/MS-DOS
COMPATION: PC-DOS/MS-DOS
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                                                                                                                                                                        MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Patent No.
                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/227,360

FILING DATE: April 13, 1994

ATTORNEY/AGENT INFORMATION:

NAME: Lech, Karen F.

REGISTRATION NUMBER: 35,238

REFERENCE/POCKET NUMBER: 00786/2

TELECOMMUNICATION INFORMATION:

TELEPHONE: (617) 542-5070
                                                                                                                                                                                                                                                                                                    TELEFAX: (617) 542-890
TELEX: 100254
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release*#1.0, Version #1.30B
CURRENT APPLICATION DATA;
APPLICATION NUMBER: US/08/310,912A
FILING DATE: September 22, 1994
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 4 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                            TYPE: amino acid
STRANDEDNESS: not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  199, Application US/08310912A
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2; Conser
                                                                                              Similarity 2; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Yu, Guo-Liang
VENTION: RPS2 GENE FAMILY, PRIMERS, PROBES, AND DETECTION
VENTION: METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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Brent, Andrew F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ausubel, Frederick M.
                                                                                                                                                                                                                                                                                                                                            (617) 542-8906
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Mindrinos, Michael N.
                                                                                                                                                                                        protein
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                                                                                                                86.7%;
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66.7%;
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                                                                                            Score 13; DB 2; : Pred. No. 3.8e+05; 0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                     00786/254001
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Pred. No. 3.8e+05;
0; Mismatches 1;
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                                                                                                                                  Length 4;
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PCT-US93-01669-41
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US-09-301-085-199
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 208
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No. 6262248
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Vea
PRICATION DATA:
APPLICATION DATA:
APPLICATION INMBER: PCT/US93/01669
FILING DATE: 01-MAR-1993
CLASSIFICATION DATA:
APPLICATION NUMBER: US/07/844,852
APPLICATION NUMBER: US/07/844,852
ATTORNEY/AGENT INFORMATION:
NAME: Wetherell, JT: Ph.D., John R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: 00786/254002
CURRENT APPLICATION NUMBER: US/09/301,085
CURRENT FILING DATE: 1999-04-28
EARLIER APPLICATION NUMBER: 08/310,912
EARLIER FILING DATE: 1994-09-22
EARLIER APPLICATION NUMBER: 08/227,360
EARLIER PLING DATE: 1994-04-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Ausubel, Frederick M
APPLICANT: Staskawicz, Brian J.
APPLICANT: Brent, Andrew F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TILE OF INVENTION: RPSZ GENE FAMILY, PRIMERS, PROBES, TILE OF INVENTION: DETECTION METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PPLICANT: Mindrinos, Michael N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Trowbridge, Ian S. APPLICANT: Collawn, Jr., Jame APPLICANT: Tainer, John A.
                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: RECEPTOR INTERNALIZATION SIGNALS NUMBER OF SEQUENCES: 57
NAME: Wetherell, Jr. Ph.D., John R. REGISTRATION NUMBER: 31,678
REFERENCE/DOCKET NUMBER: PD-1636
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
les 2; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kunkel, Barbara N
                                                                                                                                                                                                                                                                                                                                            California
                                                                                                                                                                                                                                                                                                                                                                              E: Spensley Horn Jubas & Lubitz
1880 Century Park East, Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/09301085
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Pred. No. 3.8e+09
0; Mismatches
                                                                                                                                                                                                            Version
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GENERAL INFORMA
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                                              Query Match
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TELEPAX: (619) 455-5110
INFORMATION FOR SEQ ID NO: 41:
                                                                                                                                                                                                     TELEFAX: (617) 542-8906
TELEX: 100254
INFORMATION FOR SEQ ID NO: 199:
                                                                                               STRANDEDNESS:
TOPOLOGY: lin
MOLECULE TYPE:
                                                                                                                                                                                                                             NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00786/230001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
TELEFAX: (617) 542-8906
TELEFAX: 1000-6
                                                                                                                                                                                                                                                                                                                                                                                                                     CUMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30B
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/U895/04589
FILING DATE:
CLASSTETCH
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APPLICANT:
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LENGTH: 4 amino acids
                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 4 amino acids
                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/227,360
FILING DATE: 13-APR-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
ZIP: 02110-2904
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: unknown MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Kunkel, Barbara N.
APPLICANT: Mindrinos, Michael N.
APPLICANT: Yu, Guo-Liang
TITLE OF INVENTION: RPS2 GENE AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
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            Local Similarity hes 2; Conserv
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CITY: E
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STREET: 225 Franklin Street Suite 3100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 GXY 3
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amino acid
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)GY: unknown
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RMATION:
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Katagiri, Fumiaki
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                                                                                                                    linear
                                                                                                 protein
                                                                                                                                  not relevant
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                               86.7%;
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           Score 13; DB 5;
Pred. No. 3.8e+05;
0; Mismatches 1
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Pred. No. 3.8e+05;
                                              Length 4;
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US-08-292-968-30
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US-07-962-569A-6
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APPLICANT: MENON, RAVI S.
APPLICANT: CHANG, YING-FON
APPLICANT: CHANG, YING-FON
APPLICANT: HAM, RICHARD G.
TITLE OF INVENTION: HUMAN K-CASEI
NUMBER OF SEQUENCES: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/07/962,569A
FILING DATE: 19921013
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: PEPPER PH.D., FREDERICK W.
REGISTRATION NUMBER: 31.286
REFERENCE/DOCKET NUMBER: 302224.01
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 451-1110
                                                                                                                                                                                                   Sequence 30, Application US/08292968
Patent No. 5856122
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                       uery Match
Best Local Similarity 66.7%;
Matches 2; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (619) 451-9628
INFORMATION FOR SEQ ID NO: (
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                   APPLICANT:
APPLICANT:
           TITLE OF INVENTION: MODIFICATION OF PERTUSSIS TOXIN NUMBER OF SEQUENCES: 46
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: FREDERICK W. PEPPER, PH.D.
STREET: 11545 W. BERNARDO COURT, STE. 302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: protein
                                                                                                                    APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 5 amino acids
TYPE: AMINO ACID
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                                              KLEIN, Michel H.
ARMSTRONG, Glen D.
HAZES, Bart
WYENTION
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                                                                                                                    STEIN, Penelope E. COCKLE, Stephen A. COMEN, Raymond P. LOOSMORE, Sheena
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        USA
Sim & McBurney
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Pred. No. 3.8e+05;
0; Mismatches 1
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Length 5; Indels

0,

Gaps

STREET: 5...
CITY: Toronto
Chtaric

Suite 701,

330 University Avenue

Ontario

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Sequence 36, Application US/08292968
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Best Local (
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APPLICATION NUMBER: US 08/110,947

FILING DATE: 24-AUG-1993

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/251,121

FILING DATE: 31-MAY-1994

ATTORNEY/AGENT INFORMATION:

NAME: STEWART, Michael I.

REGISTRATION NUMBER: 24,973

REFERENCE/DOCKET NUMBER: 1038-388

TELECOMMUNICATION INFORMATION:

TELEPHONE: (416) 595-1155
                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: READ, Randy J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO:
         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                         APPLICANT:
APPLICANT:
TITLE OF IN
                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
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ZIP: M5G 177
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: LEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                             STREET: Suite
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/292,968 FILING DATE:
                                                                                                                                                                                                                        ADDRESSEE: Sim & McBurney
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                No.
                                                                                                                               RY: Canada
M5G 1R7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 2; Conserv
                                                                                                                                                                                                                                                                             T: HAZES, Bart
                                                                                                                                                                 Ontario
                                                                                                                                                                                            Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                            STEIN, Penelope E. COCKLE, Stephen A. COMEN, Raymond P. LOOSMORE, Sheena KLEIN, Michel H.
                                                                                                                                                                                                                                                                                                              ARMSTRONG, Glen D.
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                                                                                                                                                                                                                                                                           MODIFICATION OF PERTUSSIS TOXIN
US/08/292,968
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Pred. No. 3.8e+05;
0; Mismatches 1
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TRANDEDNESS:

single

: 5 amino acid residues amino acid

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TELEFAX: 512-474-7577
TELEX: NOT APPLICABLE
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acid res
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 3, Application US/08618696
Patent No. 5861475
GENERAL INFORMATION:
                                                                                                     COMPUTER: IBM PC COMPATIBLE
COMPUTER: WORDERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/618,696
FILING DATE: 20-MAR-1996
CLASSIFICATION DATA:
APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION MUMBER: 07/995,269
FILING DATE: 12/21/92
ATTORNEY/AGENT INFORMATION:
NAME: PARKER, DAVID L.
REGISTRATION NUMBER: 32,165
REFERENCE/DOCKET NUMBER: UOAB:002/PAR
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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Matches 2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (416) 595-1155
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
ZIP: 77210
COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD, WHITE & DURKEE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: COOPER, Jr., J. ALLEN D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: INHIBITION OF PHAGOCYTES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 31-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Michael I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 0 FILING DATE: 24-AUG-1993 PRIOR APPLICATION DATA: APPLICATION NUMBER: US 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: HOUSTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       E: ARNOLD, WH
P.O. BOX 4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                  FLOPPY DISK
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66.7%;
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Pred. No. 3.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8e+05;
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TYPE: amino acid residues
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
US-08-618-696-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 13
US-08-618-696-14
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                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 2; Conserva
                                                                                                                                                                            RESULT 14
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MEDIUM TYPE: FLOPPY DISK
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/618,696
FILING DATE: 20-MAR-1996
CLASSIFICATION NUMBER: 07/995,269
FILING DATE: 12/21/92
ATTORNEY/AGENT INFORMATION:
NAME: PARKER, DAVID L.
REGISTRATION NUMBER: 32,165
REGISTRATION NUMBER: 32,165
REGISTRATION NUMBER: US/08/618,000
TELEPONUMINICATION INFORMATION:
TELEPONUMINICATION INFORMATION
TELEPONUMINICAT
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Best Local S
Matches 2
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Patent No. 5861475
                                                                                                                      Sequence 30,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: COOPER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: COOPER, Jr., J. ALLEN D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: INHIBITION OF PHAGOCYTES
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
ADDRESSE: ARNOLD, WHITE & DURKEE
STREET: P.O. BOX 4433
CITY: HOUSTON
STREET: HOUSTON
                                                                                                 tent No.
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                                                                           INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 GXY 3
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                                                                                                    0, Application US/08467974
5965385
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READ, Randy J.
STEIN, Penelope E.
COCKLE, Stephen A.
                                                                                                                                                                                                                                                                                                                                                         Conservative
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Pred. No. 3.8e+05;
0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                  Score 13; DB 2;
Pred. No. 3.8e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                             Length 5;
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Best Local Similarity
Matches 2; Conserve
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                                                                                                                                                                                                                                                                                                                                         RESULT 15
US-08-467-974-36
                                                                                                                                                                                                                                Patent No. 5965385
GENERAL INFORMATION:
APPLICANT: READ,
                                                                                                                                                                                                                                                                                                              Sequence 36, Application US/08467974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 24-AUG-1993
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Michael I.
REGISTRATION UNMER: 24,973
REFERENCE/DOCKET NUMBER: 1038
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/467,974
PILING DATE:
CLASSIFICATION: 435
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/467,536
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/292,968
FILING DATE: 2-AUG-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/251,121
APPLICATION NUMBER: US 08/251,121
APPLICATION NUMBER: US 08/251,121
APPLICATION NUMBER: US 08/251,121
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/110,947
APPLICATION NUMBER: US 08/110,947
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (416) 595-11:
TELEPAX: (416) 595-1163
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: Canada
ZIP: M5G 177
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: LEM PC compatible
COMPUTER: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: OOMEN, Raymond P.
APPLICANT: LOOSMORE, Sheena
APPLICANT: KLEIN, Michel H.
APPLICANT: ARMSTRONG, Glen D.
APPLICANT: HARES, Bart
APPLICANT: HARES, BOTTITLE OF INVESTION: MODIFICATION OF PERTUSSIS TOXIN
NUMBER OF SEQUENCES: 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
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                                                                                                                                                 APPLICANT: STEIN, Penelope E.
APPLICANT: COCKLE, Stephen A.
APPLICANT: COMBN, Raymond P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: Suite
CITY: Toronto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 5 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DDRESSEE:
          OF SEQUENCES:
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                                     INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       E: Sim & McBurney
Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
             HAZES, Bart
FRNTION: MODIFICATION OF PERTUSSIS TOXIN
                                                                                                                                    LOOSMORE,
                                                                                 KLEIN, Michel H.
ARMSTRONG, Glen D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       86.7%;
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Pred. No. 3.8e+05;
0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 5;
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CITY: TOTOLLO
STATE: ORGATIO
COUNTER: ORGATIO
COUNTER: Canada
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release:#1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,974
FILING DATE: 05-JUN-1995
APPLICATION NUMBER: US 08/292,968
FILING DATE: 22-AUG-1994
FRICH APPLICATION NUMBER: US 08/292,968
FILING DATE: 22-AUG-1994
PRIOR APPLICATION NUMBER: US 08/251,121
FILING DATE: 22-AUG-1994
PRIOR APPLICATION NUMBER: US 08/251,121
FILING DATE: 24-AUG-1994
PRIOR APPLICATION NUMBER: US 08/110,947
FILING DATE: 24-AUG-1993
ATTORNEY/AGENT INFORMATION:
NAME: STEMART, Michael I.
REGISTRATION NUMBER: 1038-454 MIS:vg
FILLEPAN: (416) 595-1155
TELEPAN: (416) 595-1155
INFORMATION FOR SEQ ID NO: 36:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acids
TOPOLOGY: linear
Search completed: December 13, 2004, 22:34:51 Job time: 21.4 secs
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                                                                                                                                                                      Query Match 86.7%;
Best Local Similarity 66.7%;
Matches 2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSES: Sim & McBurney
STREET: Suite 701, 330 University Avenue
CITY: Toronto
STATE: Ontario
                                                                                                                                   1 GXY 3
                                                                                        2 GTÝ 4
                                                                                                                                                                 0
                                                                                                                                                                      Score 13; DB 2; Length 5; Pred. No. 3.8e+05; 0; Mismatches 1; Indels
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Title:
Perfect score:
Sequence:
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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             1111
3210
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                                                                                                                                                                                                                                                                                               Score
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Match Length
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Gapop 10.0 ,
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15
          Published_Applications_AA:*
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Copyright (c) 1993 - 2004 Compugen Ltd
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/cgn2_6/ptodata/2/pubpaa/US09B_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US09B_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US09_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/US10A_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*
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/cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*
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/cgn2_6/ptodata/2/pubpaa/JeCT_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.pep:*
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cgn2_6/ptodata/3/m.t.
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_6/ptodata/2/pubpaa/PCTUS_PUBCOMB.pep:*
_6/ptodata/2/pubpaa/US08_NEW_PUB.pep:*
_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
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  US-10-302-617A-18
US-10-613-472-199
US-10-613-765-199
US-10-613-765-199
US-09-792-686A-34
US-10-344-779-8
US-10-344-779-8
US-10-346-779-8
US-10-394-471B-3
US-10-394-471B-3
US-10-394-471B-11
US-09-990-762-36
US-09-990-762-41
US-09-990-762-41
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Sequence 199, App
Sequence 18, Appl
Sequence 199, App
Sequence 199, App
Sequence 34, Appl
Sequence 8, Appli
Sequence 143, App
Sequence 113, Appl
Sequence 3, Appli
Sequence 11, Appl
Sequence 40, Appl
Sequence 40, Appl
Sequence 41, Appl
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equence 157,	Sequence 157, App	equence 30	equence 22,	quence 15,	equence 97,	equence 59,	equence 157	equence 35	32	equence 18	equence 12:	equence 31	quence 15,	equence 14	equence 14	equence 73:	equence 21	equence 20	equence 19	quence 18	equence 17	equence 12:	equence 10	equence 63	equence 23:	equence 23	equence 23	equence 41	equence 40	quenc	equence 6,

ALIGNMENTS

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RESULT 1

RESULT 1

RESTLA INFORMATION US/09867852

Patent No. US20020147324A1

GENERAL INFORMATION:
APPLICANT: AUSUBEL, Frederick M.
APPLICANT: Staskawicz, Brian J.
APPLICANT: Mindrinos, Michael N.
APPLICANT: Kategiri, Fundaki
APPLICANT: Kategiri, Fundaki
APPLICANT: Windrinos, Michael N.
APPLICANTION NUMBER: US/09/867,852
CURRENT FALING DATE: LARLIER FILING DATE: 1999-04-28
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/301,085
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 08/310,912
PRIOR APPLICATION DATE: 1994-04-28
PRIOR APPLICATION NUMBER: EARLIER FILING DATE: 1994-09-22
PRIOR APPLICATION DATE: 1994-09-22
PRIOR APPLICATION DATE: 1994-09-28
PRIOR APPLICATION DATE: 1994-09-27,360
PRIOR APPLICATION DATE: 1994-09-27,360
PRIOR APPLICATION DATE: 1994-04-13
NUMBER OF SEQ ID NOS: 208
SOPTWARE: FARLIER FILING DATE: 1994-04-13
NUMBER: OF SEQ ID NOS: 208
SOPTWARE: FARLIER FILING DATE: 1994-04-13
NUMBER: OF SEQ ID NOS: 208
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NUMBER: OF SEQ ID NOS: 208
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NUMBER: OF SERIER PAIL NOS: 208
SOPTWARE: FARLIER FILING DATE: 1994-04-13
NUMBER: OF SERIER PAIL N
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APPLICANT: KUNKEI, BARDATA N.
APPLICANT: Yunkei, Michael N.
APPLICANT: Yu, Guo-Liang
TITLE OF INVENTION: RPS2 GENE FAMILY, PRIMERS, PROBES, AND
TITLE OF INVENTION: DETECTION METHODS
FILE REFERENCE: 00786/254004
CURRENT APPLICATION NUMBER: US/10/613,472
CURRENT FILING DATE: 2003-07-02
PRIOR APPLICATION NUMBER: US 09/867,852
PRIOR APPLICATION NUMBER: US 09/301,085
PRIOR APPLICATION NUMBER: US 08/310,912
PRIOR APPLICATION NUMBER: US 08/310,912
PRIOR APPLICATION NUMBER: US 08/310,912
PRIOR APPLICATION NUMBER: US 08/327,360
PRIOR FILING DATE: 1994-09-22
PRIOR APPLICATION NUMBER: US 08/227,360
PRIOR FILING DATE: 1994-04-13
PRIOR FILING DATE: 1994-04-13
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 199
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Best Local Similarity 66.7
Matches 2; Conservative
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                                          Query Match
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CURRENT FILING DATE: 2002-11-22
PRIOR APPLICATION NUMBER: 60/334,434
PRIOR FILING DATE: 2001-11-30
NUMBER OF SEQ ID NOS: 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: ROZZELLE, James
APPLICANT: BOICHAKOVA, Blena
TITLE OF INVENTION: THERMUS BROCKIANUS NUCLEIC ACID POLYMERASES
FILB REFERENCE: 4768US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Ausubel, Frederick M.
APPLICANT: Staskawicz, Brian J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT CREAMISM: Thermus brockianus
                                                                                                   ORGANISM: Arabidopsis thaliana
                                                                                                                                             LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
les 2; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dahlbeck, Douglas
Katagiri, Fumiaki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Application US/10613472 o. US20040088756A1
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ilarity 66.7%;
Conservative
                   86.7%;
Score 13; DB 15;
Pred. No. 1.4e+06;
O; Mismatches 1;
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                                     Length 4;
  Indels
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US-10-613-765-199
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                                                                                              ; ORGANISM: Homo sapiens US-09-792-686A-34
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CURRENT FILING DATE: 2003-07-02
PRIOR APPLICATION NUMBER: US 09/867,852
PRIOR FILING DATE: 2001-05-29
PRIOR APPLICATION NUMBER: US 09/301,085
PRIOR PILING DATE: 1999-04-28
PRIOR APPLICATION NUMBER: US 08/310,912
PRIOR APPLICATION NUMBER: US 08/310,912
PRIOR APPLICATION NUMBER: US 08/227,360
PRIOR APPLICATION NUMBER: US 08/227,360
Query Match
Best Local Similarity bb...
Conservative
                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Goronzy, Jorg J.
APPLICANT: Kopecky, Stephen L.
APPLICANT: Weyand, Cornelia M.
TITLE OF INVENTION: METHODS AND MATERIALS FOR EVALUATING
TITLE OF INVENTION: CARDIOVASCULAR CONDITIONS
                                                                                                                                                              NUMBER OF SEQ ID NOS: 37
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 34
                                                                                                                                                                                                                                                                                                                                                                              Sequence 34, Application US/09792686A Publication No. US20030068645A1
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                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/792,686A CURRENT FILING DATE: 2001-02-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR FILING DATE: 1994-04-13
NUMBER OF SEQ ID NOS: 214
SOFTWARE: FastSEQ for Windows Version 4.0
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APPLICANT: SALMERON, JOHN
TITLE OF INVENTION: RPSZ GENE FAMILY, PRIMERS,
TITLE OF INVENTION: DETECTION METHODS
FILE REFERENCE: 00786/254005
                                                                                                                                                                                                                                                         FILE REFERENCE: 07039-250001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 4
TYPE: PRT
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                                                                                                                                  LENGTH: 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STASKAWICZ, BALLESTASKAWICZ, BALLESTARI, FUMIAKI
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5. US20040172673A1
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                                         86.7%;
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                            0; Mismatches
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                                           Score 13; DB 10;
Pred. No. 1.4e+06;
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                                                            Length 5;
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FILE REFERENCE: GP50033

CURRENT APPLICATION NUMBER: US/10/344,779

CURRENT FILING DATE: 2003-02-17

PRIOR APPLICATION NUMBER: PCT/US01/26161

PRIOR PILING DATE: 2001-08-21

PRIOR PILING DATE: 2000-08-21

PRIOR FILING DATE: 2000-08-21

PRIOR FILING DATE: 2000-08-21

PRIOR APPLICATION NUMBER: US 60/226,524

PRIOR PILING DATE: 2000-09-07

INUMBER OF SEQ ID NOS: 18

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 8

LENGTH: 5

TYDE: DET
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US-10-376-121A-143
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Best Local 9
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APPLICANT: WATTAM, TREVOR A.
TITLE OF INVENTION: USEFUL IN TREATMENT OF RANK LIGAND MEDIATED DISORDERS
TITLE OF INVENTION: USEFUL IN TREATMENT OF RANK LIGAND MEDIATED DISORDERS
CURRENT APPLICATION DATA: U$/10/376,121A
APPLICATION NUMBER: U$/10/376,121A
FILING DATE: 27-Mar-2003
CLASSIFICATION: <UNKnown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/867,819
FILING DATE: April 13, 1992
APPLICATION NUMBER: 07/648,205
FILING DATE: January 31, 1991
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                                                                                                                                                                                           COUNTRY: USA
ZIP: 30309-3400
COMPUTER READABLE FORM:
WEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Patrea L. Pabst
STREET: Suite 2000, 1201 West Peachtree Street, N.B.
CITY: Atlanta
STATE: GA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Harley, John
TITLE OF INVENTION: METHODS AND REAGENTS FOR DIAGNOSIS OF
AUTOANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 GXY 3
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Pred. No. 1.4e+06
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RESULT 9 US-10-394-471B-3

Sequence 3, Application US/10394471B Publication No. US20040185047A1 GENERAL INFORMATION:

Jill; Carton, Jill; Scallon, Bernard

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APPLICANT: TRUNCH, ALENSEGED
APPLICANT: WATTAM, TREVOR A.
FITTLE OF INVENTION: ANTI-RANK LIGAND MONOCLONAL ANTIBODIES
FILE OF INVENTION: USEFUL IN TREATMENT OF RANK LIGAND MEDIATED DISORDERS
FILE REFERENCE: G95033
CURRENT APPLICATION NUMBER: US/10/738,809
CURRENT APPLICATION NUMBER: US/10/344,779
PRIOR APPLICATION NUMBER: US/10/344,779
PRIOR APPLICATION NUMBER: US/10/344,779
PRIOR APPLICATION NUMBER: US/10/344,779
PRIOR APPLICATION NUMBER: US/05/26161
PRIOR APPLICATION NUMBER: US/07/26161
PRIOR APPLICATION NUMBER: US/07/26161
PRIOR APPLICATION NUMBER: US/07/26161
PRIOR APPLICATION NUMBER: US/07/26161
PRIOR FILING DATE: 2000-08-21
PRIOR FILING DATE: 2000-08-21
PRIOR APPLICATION NUMBER: US/07/230,639
PRIOR APPLICATION STATE SOUND SUBJECT OF SEQ ID NOS: 18
SOFTWARE: FASTSEQ for Windows Version 4.0
B,
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                                                                           Query Match 86.7
Best Local Similarity 66.7
Matches 2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 8, Application US/10738809
Publication No. US20040171117A1
GENERAL INFORMATION:
APPLICANT: SWEET, RAYMOND W.
APPLICANT: TORNETTA, MARK A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 65...
Thes 2; Conservative
                                                                                                                                                                                               TYPE: PRT
ORGANISM: Mus musculus
                                                                                                                                                                                                                                                  LENGTH: 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 07/472,947
FILING DATE: January 31, 1990
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: peptide SEQUENCE DESCRIPTION: SEQ ID NO: 143:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Pabst, Patrea L.
REGISTRATION NUMBER: 31,2
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acid
                                         1 GXY 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (404)-817-8473
TELEFAX: (404)-817-8588
                                                                                                          86.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                86.7%; Score 13; DB 14; Length 5;
66.7%; Pred. No. 1.4e+06;
tive 0; Mismatches 1; Indels
                                                                                  Score 13; DB 16;
Pred. No. 1.4e+06;
0; Mismatches 1
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BER: OMRF114CIP(2)DIV(2)
                                                                                                                           Length 5
                                                                                     Indels
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Query Match
Best Local Similarity
Matches 2; Conserve
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US-10-394-471B-11
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                                                                                                                                                                                                                                                                                                                                            US-09-990-762-36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Giles-Komar, Jill; Carton, Jill; Scallon, Bernard J.
TITLE OF INVENTION: ANTI-TNF ANTIBODIES, COMPOSITIONS, METHODS AND USES
FILE REFERENCE: CENO202
CURRENT APPLICATION NUMBER: US/10/394,471B
CURRENT FILING DATE: 2003-03-21
PRIOR APPLICATION NUMBER: 60/367,903
PRIOR FILING DATE: 2002-03-26
NUMBER OF SEQ ID NOS: 17
SOFTWARE: Patentin Ver 3.1
SEQ ID NO 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/367,903
PRIOR FILING DATE: 2002-03-26
NUMBER OF SEQ ID NOS: 17
SOPTWARE: PATENTIN VET 3.1
SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 11, Application US/10394471B Publication No. US20040185047A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                 Sequence 36, Application US/09990762
Patent No. US20020119498A1
SOFTWARE: PatentIn Ver. SEQ ID NO 36
                                                                                                                                                                                                                                                                                    GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: CEN0202
CURRENT APPLICATION NUMBER: US/10/394,471B
CURRENT FILING DATE: 2003-03-21
                                  APPLICANT: JOUNG, J. KEITH
APPLICANT: MILLER, JEFFREY
APPLICANT: MILLER, JEFFREY
APPLICANT: MILLER, JEFFREY
APPLICANT: PABO, CCARL O,
APPLICANT: PABO, CARL O,
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INTERACTION TRAP ASSAYS
FILE REFERENCE: MTV-030.02 (20021-3002)
CURRENT APPLICATION NUMBER: US/09/990,762
CURRENT FILING DATE: 2001-11-14
PRIOR APPLICATION NUMBER: 09/858,852
PRIOR APPLICATION NUMBER: 60/204,509
PRIOR FILING DATE: 2001-05-16
PRIOR FILING DATE: 2000-05-16
NUMBER OF SEQ ID NOS: 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: ANTI-THE ANTIBODIES, COMPOSITIONS, METHODS AND USES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT ORGANISM: Homo sapiens
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66.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 13; DB 16;
Pred. No. 1.4e+06;
0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .4e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 5;
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; OTHER INFORMATION: Description of Artificial Sequence: Zinc finger; OTHER INFORMATION: recognition sequence US-09-990-762-36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 13
US-09-990-762-41
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 41, Application US/09990762
Patent No. US20020119498A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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FILE REFERENCE: MTV-030.02 (20021-3002)
CURRENT APPLICATION NUMBER: US/09/990,762
CURRENT FILING DATE: 2001-11-14
PRIOR APPLICATION NUMBER: 09/858,852
PRIOR FILING DATE: 2001-05-16
PRIOR FILING DATE: 2001-05-16
PRIOR FILING DATE: 2000-05-16
NUMBER: 09/204,509
PRIOR FILING DATE: 2000-05-16
NUMBER: 09 SEQ ID NOS: 97
                                                                                                                      APPLICANT: JOUNG, J. KEITH
APPLICANT: MILLER, JEFFREY
APPLICANT: MILLER, JEFFREY
APPLICANT: PABO, CARL O.
APPLICANT: PABO, CARL O.
ITIE OF INVENTION: METHODS AND COMPOSITIONS FOR INTERACTION TRAP ASSAYS
FILE REFERENCE: MTV-030.02 (20021-3002)
FIL
                                                                                                    SEQ ID NO 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: JOUNG,
APPLICANT: MILLE
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ORGANISM: Artificial Sequence
                                                           ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ENGTH:
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2; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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Pred. No. 1.4e
0; Mismatches
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Pred. No. 1.4e+06;
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Sequence 6, Application US/09518737

Publication No. US20030008321A1

GENERAL INFORMATION:
APPLICANT: FUXUI, YASUHISA
APPLICANT: WAGATA, SATOSHI
APPLICANT: NAGATA, SATOSHI
APPLICANT: SAITO, NAOAKI
TITLE OF INVENTION: MONOCLONAL ANTIBODY RECOGNIZING
TITLE OF INVENTION: HOSPHATIDYLINOSITOL-3,4-DIPHOSPHATE
FILE REFERENCE: 1965/49618
CURRENT APPLICATION NUMBER: US/09/518,737
CURRENT FILING DATE: 2000-03-03
PRIOR APPLICATION NUMBER: UB 1999-250209
PRIOR FILING DATE: 1999-09-03

NUMBER OF SEQ ID NOS: 10
TYPE: PRT

ORGANISM: Artificial Sequence

FEMTURE:

OTHER INFORMATION: Description of Artificial Sequence: Zinc finger

OTHER INFORMATION: recognition sequence

US-09-858-852A-36
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                                                                                                                  APPLICANT: JOUNG, J. KEITH
APPLICANT: JULER, JEFFREY
APPLICANT: MILLER, JEFFREY
APPLICANT: PABO, CARL O.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INTERACTION TRAP ASSAYS
FILE REFERENCS: MTW-030.01 (20021-3001)
CURRENT APPLICATION NUMBER: US/09/858,852A
CURRENT FILING DATE: 2001-05-16
PRIOR APPLICATION NUMBER: 60/204,509
PRIOR FILING DATE: 2000-05-16
NUMBER OF SEQ ID NOS: 91
SOCTWARE: PATENTIN VET. 2.1
SEQ ID NO 36
LENGTH: 6
LENGTH: 6
LENGTH: 6
                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 36, Application US/09858852A Publication No. US20030044787A1 GENERAL INFORMATION:
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Best Local Similarity
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SEQ ID NO 6
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ORGANISM: Mus musculus
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Local Similarity 66.7%;
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Query Match

Score 13;

DB 10; Length 6;

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Best Local Similarity 66.7%; Pred. No. 1.4e+06; Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps

Qy 1 GXY 3

Db 3 GAY 5
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Search completed: December 13, Job time : 69.6 secs

2004, 23:02:40

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Sequence:
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Maximum DB seq length: 2000000000
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Maximum Match 100%
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Copyright (c) 1993 - 2004 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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alkaline serine pr	alphaA-conotoxin P	guanidino kinase M	cysteine proteinas	fibrinogen gamma c	T-cell receptor be	seed protein ws-9	alpha 1-proteinase	Ig heavy chain V r	H+-transporting tw	tyrocidine synthas	Ig heavy chain V r	T-cell receptor J-	T-cell receptor J-	Ig heavy chain CDR	pyrrhocoricin - Py

ALIGNMENTS

Query Match 86.7%; Score 13; DB 2; Length 9; Best Local Similarity 66.7%; Pred. No. 2.8e+05; Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0; Qy 1 XY 3	RESULT 2 G41946 T-cell receptor gamma chain (2t.23) - mouse (fragment) C.Species: Mus musculus (house mouse) C.JDate: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 07-May-1999 C.Accession: G41946 R.Mosley, R.L.; Whetsell, L.; Schaefer, F.V.; Miller, K.S.; Klein, J.R. Mol. Cell. Biol. 11, 5902-5909, 1991 A;Title: Rearrangement and junctional-site sequence analyses of T-cell receptor gamma g A,Reference number: A41946; MUID:92049316; PMID:1658619 A;Status: preliminary; not compared with conceptual translation A,Molecule type: DNA A,Residues: 1-9 < WHE> C;Keywords: T-cell receptor	RESULT 1 S36550 Ig heavy chain V region - mouse C;Species: Mus musculus (house mouse) C;Apte: 06-Jan-1994 #sequence_revision 01-Dec-1995 #text_change 05-Nov-1999 C;Accession: 33650 R;Jacob, J.; Kelsoe, G. submitted to the EMBL Data Library, July 1992 A;Description: In situ studies on the primary immune response to (4-hydroxy-3-nitrophen A;Reference number: S25024 A;Reference number: S25024 A;Rocession: S36850 A,Status: preliminary A;Accession: S36850 A,Cross-references: EMBL:X67387; NID:g50113; PIDN:CAA47799.1; PID:e51594; PID:g1333871 C;Keywords: heterotetramer; immunoglobulin Query Match Best Local Similarity 66.7%; Score 13; DB 2; Length 9; Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0; Qy 1 GXY 3 Db 6 GSY 8

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C; Accessic
R; Yamada,
RESULT
S47391
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C;Species: Plasmodium falciparum
C;Date: 24-Aug-1990 #sequence_revision 24-Aug-1990 #
C;Accession: A33099
R;Nichols, J.H.; Hager, L.P.
submitted to the Protein Sequence Database, May 1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G. J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT0250
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A; Residues: 1-11 <NAK>
C; Keywords: T-cell receptor
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J. Exp. Med. 173, 1091-1097, 1991
A;Title: T cell receptor V gene usage of islet beta cell-reactive T
A;Reference number: PT0209; MUID:91217621; PMID:1902501
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A; Residues: 1-11 < YAM>
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C; Keywords: heterotetramer;
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A;Accession: A33099
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             y heavy chain CRD3 region (clone 2-109B) - human (fragment)
;Species: Homo sapiens (man)
;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
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0; Mismatches 1
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R;Lehner, P.J.
submitted to the EMBL Data Library, August 1994
A;Description: Human HLA-A0201 restricted recogn
A;Reference number: $47355
A;Accession: $47391
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R;Lehner, P.J.
submitted to the EMBL Data Library, August 1994
A;Description: Human HLA-A0201 restricted recognition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: EMBL:Z35712; NID:g527519; PIDN:CAA84781.1; C;Keywords: T-cell receptor
                                                                                                  A; Experiment C; Keywords:
                                                                                                                                     A; Molecule type: DNA
A; Residues: 1-12 <LEV>
                                                                                                                                                                      A;Title: Molecular characterization of transgene-induced A;Reference number: PH1580; MUID:93301609; PMID:8315387 A;Accession: PH1611
                                                                                                                                                                                                                         R;Levinson, D.A.; Campos-Torres, J.;
J. Exp. Med. 178, 317-329, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: EMBL:23
C;Keywords: T-cell receptor
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A;Residues: 1-12 <LEH>
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A;Residues: 1-12 <LEH>
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                                                                                                                                                                                                                                                      g H chain V-D-J region (wild-type clone 341) - mouse ($pecies: Mus musculus (house mouse)
Date: 02-Jun-1994 #sequence revision 02-Jun-1994 #te:
                                                                                                  Experimental source: bone marrow; Keywords: immunoglobulin
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Pred. No. 8.8e+02;
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Pred. No. 8.8e+02;
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A;Reference number: $473
A;Accession: $4736
A;Accus: preliminary
A;Molecule type: mRNA
A;Residues: 1-13 <LEH>
                                                                                                                T-cell antigen receptor VJ junction beta chain - human ()Species: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change C;Accession: $47368
R;Lehner; P,J.
submitted to the EMBL Data Library, August 1994
A;Description: Human HIA-A0201 restricted recognition of influen A;Reference number: $47355
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A;Reference number: PH0746; MUID:92078846; PMID:1836010
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C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PH0802
R;Casanova, J.L; Romero, P.; Widmann, C.; Kourilsky, P.; Maryanski, J.L.
J. Exp. Med. 174, 1371-1383, 1991
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J. Exp. Med. 174, 1371-1383, 1991
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A; Residues: 1-12 < CAS>
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allelic exclusion and antigen-specific repertoire.
A;Reference number: PH0746; MUID:92078846; PMID:1836010
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Matches 2
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Cross-references: EMBL:Z35693; NID:g527477; PIDN:CAA84762.1; PID:g527478
Keywords: T-cell receptor
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Experimental source: T lymphocyte
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Residues: 1-12 <CAS>
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pred. No. 8.8e+02;
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Pred. No. 8.8e+02;
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unidentified 6.3/40K protein [imported] - rice (;Species: Oryza sativa (rice) (;Date: 20-Apr-2000 #sequence_revision 20-Apr-2) (;Accession: PQ0700 #sq.Komatsu, S.; Kajiwara, H.; Hirano, H.
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PH0788
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C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PH0788
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A;Molecule type: mRNA
A;Residues: 1-13 < LEH>
A;Cross-references: EMBL:Z3:
C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                             A;Experimental source: T lymphocyte C;Keywords: T-cell receptor
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A; Residues: 1-13 < CAS>
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allelic exclusion and antigen-specific repertoire.
A;Reference number: PH0746; MUID:92078846; PMID:1836010
A;Accession: PH0788
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J. Exp. Med. 174, 1371-1383,
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A; Accession: $47383
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A;Description: Human HLA-A0201 restricted recognition of
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Appl. Genet. 86, 935-942, 1993 e: A rice protein library; a data-
rence number: PQ0696
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Pred. No. 9.5e+02;
0; Mismatches 1;
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RESULT 15
A59018
NUC1 enhancer binding protein 70K chain MUC1EBP-70 - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 08-May-1998 #sequence_revision 08-May-1998 #text_change 08-May-1998
C;Accession: A59018
R;Abe, M.; Smith, C.J.; Larson, C.J.
R;Abe, M.;Abe, M.;Abe, M.;Abe, M.;Abe, M.;Abe, M.;Abe, M.
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A, Molecule type: protein
A, Residues: 1-13 - KCM's
A, Cross-references: UNIPROT: Q7M1U1
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Search completed: December 13, 2004, 22:32:59
Job time : 15.2 secs
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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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length: 2000000000
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Gapop 10.0 , Gapext 0.5
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    GenCore version
(c) 1993 - 2004
Q958E4
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Q9qwto mus musculu
Q8ney9 homo sapien
Aao63472 homo sapien
Q9uc46 homo sapien
Q7up62 homo sapien
P83109 tenebrio mo
 Aac39714 homo sapi
Aab92363 homo sapi
Q9gjw1 lagenorhync
Q9gjw2 lagenorhync
Q9gjw3 lagenorhync
Q61d78 clostridium
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Q6jdk5
Q9smv3
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5 canis famil
3 arabidopsis
9 homo sapien
5 saguinus oe
4 homo sapien
6 spinacia ol
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1 torpedo mar
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Q9S8H8	Q9GNL7	Q25281	PYRR_PYRAP	Q9F6J8	Q9R4F8	Q47895	Q9GNL6	Q9N6I3	AAB20893	Q9PRX1	Q9R501	Q9R4F9	Q9R4C6
Q9s8h8	Q9gnl7	Q25281	P37362	Q9£6j8	Q9r4f8	Q47895	Q9gn16	Q9n6i3	Aab208	Q9prx1	Q9r501	Q9r4f9	Q9r4c6
brassica na	drosophila	leishmania	pyrrhocoris	yersinia pe	aeromonas.	fremyella d		toxoplasma	Aab20893 clostridi	struthio ca	bacillus su	aeromonas.	agrobacteri

ALIGNMENTS

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RESULT 1

UF06_MOUSE STANDARD; PRT; 8 AA.

ID UF06_MOUSE STANDARD; PRT; 8 AA.

AC P38644;

DT 01-OCT-1994 (Rel. 30, Created)

DT 01-OCT-1994 (Rel. 30, Last sequence update)

DT 01-UT1-2004 (Rel. 44, Last annotation update)

DT 05-UT1-2004 (Rel. 44, Last annotation update)

DE Unknown protein from 2D-PAGE of fibroblasts (P50) (Fragment)

OS Mus musculus (Mouse).
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Best Local S
Matches 2
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Q9QWT0;
01-MAY-2000
01-MAY-2000
01-DEC-2001
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NON TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE=Fibroblast;

MEDLINE=95009907; PubMed=7523108;

Merrick B.A., Patterson R.M., Wichter L.L., He C., Selki "Separation and sequencing of familiar and novel murine preparative two-dimensional gel electrophoresis.";

Electrophoresis 15:735-745(1994)

-I-MISCELLANEOUS: On the 2D-gel the determined pl of the protein is: 5.2, its MW is: 50 kDa.
STRAIN-Sv129 D3;
Lutz P.G., Moog-Lutz C., Houzel-Charavel A., C
Submitted (JUN-1998) to the EMBL/GenBank/DDBJ
EMBL; AJ007030; CAA07429.1; -.
                                                                                                             Eukaryota; Metazoa;
Mammalia; Eutheria;
NCBI_TaxID=10090;
                                                                                                                                                                                               Proteinase 3
Name=Prtn3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                              Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=10090;
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O (TrEMBLrel. 13,
1 (TrEMBLrel. 19,
3 (Myeloblastin)
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Rodentia;
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Last annotation update)
(Fragment).
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SEQUENCE
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NON_TER
SEQUENCE
                                                                                                  SEQUENCE FROM N.A.
Bairwa N.K., Bamezai R.;
Bairwa N.K., Bamezai R.;
"Study of promoter polymorphism
Submitted (DEC-2002) to the EMBL
EMBL; AY203963; AAO63472.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
05-UUL-2004 (TrEMBLrel. 27, Last annotation update)
Type II hair-specific keratin (Type II hair keratin)
Name=KRTHB6;
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NON TER 9 9
SEQUENCE 9 AA; 937 M
                                                                Keratin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bairwa N.K., Bamezai R.;
Submitted (DEC-2002) to the EME
EMBL; AY037552; AAK68688.1; -.
EMBL; AY203963; AAO63472.1; -.
GO; GO:0005882; C:intermediate
                                                                                                                                                                                                                                                   Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                02-MAR-2004 (TrEMBLrel. 27, Created)
02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
Type II hair keratin (Fragment).
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Submitted (MAY-2001)
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                                                                                                                                                                                                                                                                                                                                                                                                                      AA063472
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0; Mismatches
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EMBL/GenBank/DDBJ databases.
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RESULT 6
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O1-C
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Best Local S
Matches
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Matches 2
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Matches 2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE.

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01-OCT-2003
01-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Q9UC46

Q9UC46;
Q9UC46;
Q1-MAY-2000 (TrEMBLrel. 13, Created)
Q1-MAY-2000 (TrEMBLrel. 13, Last sequence update)
Q1-MAY-2003 (TrEMBLrel. 24, Last annotation update)
Q1-JUN-2003 (TrEMBLrel. 24, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
Liu Y., He X., Xu L.,
Submitted (JUN-2003)
                                                                                                                                                                                                                                                                                                                                                                     EMBL; AY328516; AAP94210.1;
NON_TER 1 1
                                                                                                                                                                                                                                                                                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Name=HLA-F;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (Human)
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lomo sapiens (Human)
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2; Conser
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GSY
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(TrEMBLrel. 25, Last sequence update)
(TrEMBLrel. 25, Last annotation update)
b antigen (Fragment).
                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                  1208 MW;
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Primates;
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                                                                                                                                                                 Score 13; DB Pred. No. 7.2e 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Y.;
e EMBL/GenBank/DDBJ
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No. 6.60
0; Mismatches
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Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                                    DB
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                                                                                                                                                                                                                                               Length 11;
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RESULT 7
ADFB_TENMO

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STANDOC PRESENTANT OCCUPANTA PROPERTIES OCCUPANTA P
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Best Local S
Matches 2
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PRELIMINARY; PRT; 13 AA.
P82432;
P82432;
O1-JUN-2000 (TrEMBLrel. 14, Created)
O1-JUN-2000 (TrEMBLrel. 14, Last sequence update)
O1-JUN-2000 (TrEMBLrel. 24, Last annotation update)
32 kDa cell wall protein (Fragment)
Nicotiana tabacum (Common tobacco).
Nicotiana tabacum (Common tobacco).
Sukaryota; Viridiplantae; Streptophyta; Embryophyta; T
Spermatophyta; Magnoliophyta; eudicotyledons; core eud
lamiids; Solanales; Solanaceae; Nicotiana.
(NCBI_TaxID=4097;
                                                                                                                                                                                                                                                                                                                                                                SEQUENCE.
SETRAIN-CV. PETIT HAVANA;
STRAIN-CV. PETIT HAVANA;
Blee K.A., Bonham V.A., Mitchell G.P., Ro
Wojtaszek P., Bolwell G.P.,
Wojtaszek P., Bolwell G.P.,
Thermal Study of Secondary cell wall
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADFB_TENMO STANDARD; PRT; 13 AA. P83109; P83109; Rel. 42, Created)
10-OCT-2003 (Rel. 42, Last sequence update)
10-OCT-2003 (Rel. 44, Last annotation update)
Antidiuretic factor B (ADFb).
Tenebrio molitor (Yellow mealworm).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota; Meoptera; Endopterygota; Coleoptera; Polyphaga; Cucujiformia; Tenebrionidae; Tenebrio.
                                                                                                                                                                                                                                                                                               Planta 0:0-0(2000).
-!- SUBCELLULAR LOCATION: CELL WALL.
-!- TISSUE SPECIFICITY: XYLEM:
                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                           Cell wall.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Direct protein sequencing; Hormone; Neuropeptide. SEQUENCE 13 AA; 1562 MW; 0240A4504B8A632B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SYNTHESIS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE, FUNCTION, TISSUE
                                                                                                                                                                                                                  TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CGMP as second messenger. May function as an antidiuretic hor SUBCELLULAR LOCATION: Secreted.
TISSUE SPECIFICITY: Brain. Appears to be mainly expressed in pairs of bilaterally symmetrical cells in the protocerebrum. MASS SPECTROMETRY: MW=1560.33; METHOD=MALDI; RANGE=1-13; NOTE-Ref.1.
                                                                                                                                                                                                                                                                   GO: 0005618; C:cell wall;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4
                                                             Similarity 66. 2; Conservative
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                                                                                                                                                                                AA;
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iehart U.M., Nicolson
                                                                                                                                                                                1307 MW;
                                                                                            86.7%;
66.7%;
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                                                          Score 13; DB:
Pred. No. 8.3e
0; Mismatches
                                                                                                                                                                                                                                                                   IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 13;
Pred. No.
                                                                                                                                                                             D5022297D8697DC6 CRC64;
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RESULT 11
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RESULT 10
Q7YOR1
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Q7YOR1; PRT:
01-OCT-2003 (TrEMBLrel. 25, Create:
01-OCT-2003 (TrEMBLrel. 25, Last s:
01-MAR-2004 (TrEMBLrel. 26, Last s:
01-MAR-2004 (TrEMBLrel. 26, Last s:
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Q7M1U1;
01-MAR-2004
01-MAR-2004
01-MAR-2004
                                                                                                     Ligase.
NON TER
SEQUENCE
                                                                                                                                         Wachter A., Steininger H., Rausch T., Bogs J
Submitted (MAY-2003) to the EMBL/GenBank/DDE
EMBL, AJ564376; CAD91995.1; -.
GO; GO:0004357; F:glutamate-cysteine ligase
GO; GO:0016874; F:ligase activity; IEA.
                                                                                                                                                                                                                                                  Brassica juncea (Leaf mustard) (Indian mustard).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosid
eurosids_II; Brassicales; Brassicaceae; Brassica.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "A rice protein library; a data-file two-dimensional electrophoresis."; Theor. Appl. Genet. 86:935-942(1993). POR, PC0700; PO0700.

NON_TER 13 13
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01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Unidentified 6.3/40K protein (Fragment).
Oryza sativa (Rice).
Eukaryota, Viridiplantae, Streptophyta; Embryophyta
Spermatophyta; Magnoliophyta; Liliopsida; Poales; P
Ehrhattoideae; Oryzeae; Oryzaa.
                                                                                                                                                                                                                                                                                                           Name=gsh1
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2; Conserv
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GAY 11
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to the EMBL/GenBank/DDBJ
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Last annotation update)
hetase (EC 6.3.2.2) (Fragment).
                                                Score 13; DB
Pred. No. 8.3e
0; Mismatches
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; Poales; Poaceae;
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RESULT 12
UC14 MAIZE STANDARI
ID TUC14 MAIZE STANDARI
AC P80620;
DT 01-0CT-1996 (Rel. 34, 0
DT 01-0CT-1996 (Rel. 34, 1
DT 05-UUL-2004 (Rel. 44, 1
DE Unknown protein from 21
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Best Local
  Query Match
Best Local
                                                                                                                                                    Touzet P., Riccardi F., Morin C., Damerva.
Permollet J.-C., Zivy M., de Vienne D.;
"The maize two dimensional gel protein dal
genome analysis program.";
Theor. Appl. Genet. 93:997-1005(1996).
-!- MISCELLANEOUS: On the 2D-gel the dete:
protein is: 5.9, its MW is: 34.6 kDa.
MaizeDB; 123944;
-- Prince MaizeDB; 123944;
MaizeDB; 123944;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD clade; Panicoideae; Andropogoneae; Zea.
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"Prunus serotina amygdalin hydrolase
Plant Physiol. 100:282-289(1992).
-!- CATALYTIC ACTIVITY: (R)-prunasin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids I; Rosales; Rosaceae; Amygdaloideae; Prunus.
                                                                                       NON TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (Fragment).
Zea mays (Maize)
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                                                                                                                                        Direct
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   embryonal tissues. PTM: Glycosylated.
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SUBUNIT: Monomer
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2; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (Black cherry)
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34, Last sequence update)
44, Last annotation update)
rom 2D-PAGE of etiolated col
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66.7%;
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Pred. No. 9.4e
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  DB 1;
9.4e+03;
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COC Eukar
OC Sperm
OC PACCO
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RESULT 15
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ID Q4789

AC Q4789
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01-JUN-2003 (TrEMBLrel. 24, Last annotation
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GO; GO:00008422; Fibera-glucosidase activity; IEA.
SEQUENCE 15 AA; 1667 MW; 46C4037DDC8C730B CRC64;
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01.MAY-2000 (TrEMBLrel. 13, Last sequence update)
01.UN-2003 (TrEMBLrel. 24, Last annotation update)
Beta-D-glucoside glucohydrolase (EC 3.2.1.21) (Fragment)
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Fremyella diplosiphon (Calothrix PCC Bacteria; Cyanobacteria; Nostocales;
   Q47893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.

MEDLINE=6623345; PubMed=3086870;

MEDLINE=6623345; PubMed=3086870;

Conley P.B., Lemaux P.G., Lomax T.L., Grossman A.R.;

Conley P.B., Lemaux P.G., Lomax T.L., Grossman A.R.;

"Genes encoding major light-harvesting polypeptides are

the genome of the cyanobacterium Fremyella diplosiphon.

Proc. Natl. Acad. Sci. U.S.A. 83:3924-3928(1986).
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AC 047893;
DT 01-NOV-1996 (TrEMBLrel 01, Last sequence update)
DT 01-NOV-1996 (TrEMBLrel 08, Last annotation update)
DE Beta-phycocyanin (Fragment).

OC DEACHER 197;
RN 11
RP PROPER 197;
RN 11
RP SEQUENCE FROM N.A.
RP SEQUENCE FROM N.A.
RP SEQUENCE 15 AA; Lomax T.L., Grossman A.R.;
RP "Genes encoding major light-harvesting polypeptides are clustered on the genome of the cyanobacterium Fremyella diplosiphon.";
RP Proc. Natl. Acad. Sci. U.S.A., 83:3924-3928(1986).
Proc. Natl. Acad. Sci. U.S.A., 83:3924-3928(1986).
RR SEQUENCE 15 AA; 1528 MW; 7FF2F65518F493D4 CRC64;
Query Match
Best Local Similarity 66.7%; Score 13; DB 2; Length 15;
Best Local Similarity 66.7%; Pred. No. 9.4e+03;
Matches 2; 'Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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ALIGNMENTS

RESULT 1 AAP61656

AAP61656 standard; protein; 4 AA.

25-MAR-2003 03-OCT-2002 21-AUG-1991

(revised) (revised) (first entry)

independent protein

AAP61656;

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WPI; 1986-118872/18.
                                                                                  Kettner CA,
                                                                                                                      16-NOV-1983;
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                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                       Cell growth inhibitor; tumour cell growth inhibitor,
                                                                                                                                                                                                                   Sequence of peptide which inhibits cyclic-nucleotide kinase activity and mammalian cell growth.
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The cpds. of the invention inhibit protein phosphorylation. The inventoclaim a process for inhibiting the growth of tumour cells in a medium which comprises contacting the cells with a cpd. of formula (AAP61654-P61661) or a physiologically acceptable salt. (Updated on 03-OCT-2002 the dark of the correct PA field.)

The inventors

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Claim 1; Col 4; 9pp; English.

Inhibition of tumour cell growth - using peptide and aminoacid halo: $methyl\ ketone(s)$.

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RESULT 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This peptide is useful as part of a peptide/halo-methyl ketone cpd., for treating picornavirus, egpolio- or rhinovirus infections. It inhibits the processing of picornavirus capsid proteins by virus encoded proteases. See AAP71301-12. See also US4652552. (Updated on 25-MAR-2003 to correct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Treating picorna-virus infection with peptide halo: methyl ketone cpds. esp. for treating polio virus and rhino virus infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Picornaviridae; poliovirus; rhinovirus; antiviral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Peptide component of cpd. for treating picornavirus infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-MAR-2003
15-MAY-1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 4 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 4; 10pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kettner CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-AUG-1984;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-AUG-1984;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-JAN-1987
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                                                                           25-MAR-2003
30-JAN-1995
                                                                                                                                                  AAR55333;
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                                                                                                                                                                                             AAR55333 standard; peptide; 4 AA.
                           Inhibitor of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N
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                                                                                                                                                                                                                                                                                                                                                                                                        2; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GAY 4
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                                                                                                                                                                                                                                                                                                                  GAY 4
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(first en
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                           tyrosine kinase
                                                                           (revised)
(first entry)
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Pred. No. 1.7e+06;
0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 13; DB 1;
Pred. No. 1.7e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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RESULT 4
AAR75499
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                    The sequence is that of a protein tyrosine kinase inhibitor contg. D- Tyr or tetrafluoro-Tyr residues. The peptide can be used to inhibit TK-mediated signal transduction and is useful for controlling proliferative diseases, e.g. cancer, psoriasis and to treat viral, inflammatory, allergic and cardiovascular diseases. See also AAR55302-39. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New peptide inhibitors of protein tyrosine kinase - contain D-Tyr or tetrafluoro-Tyr residue, for treating proliferative diseases, viral infection, inflammation etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-NOV-1992;
27-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PTK; signal transduction; proliferation; cancer; psoriasis; viral;
inflammatory; allergic; cardiovascular; diseases.
                                                                                                                                                                                                                                                                                                                                                                Sequence 4 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 7; Page 36; 44pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1994-183419/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dobrusin
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                                                                                                                                   CSa;
                                                                                                                                                                          25-MAR-2003
13-NOV-1995
                                                                                                                                                     C5a receptor-antagonist core tetrapeptide
                                                                                                                                                                                                         AAR75499;
                                                                                                                                                                                                                             AAR75499 standard; peptide; 4 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (WARN ) WARNER LAMBERT CO.
                               06-DEC-1993;
                                                  16-NOV-1994;
                                                                      15-JUN-1995.
                                                                                          WO9516033-A1
                                                                                                              Homo sapiens
           (CIBA ) CIBA
                                                                                                                                 C5a receptor-antagonist; antiinflammatory
                                                                                                                                                                                                                                                                                                     1 GXY 3
                                                                                                                                                                                                                                                                                                                       Similarity
2; Conserv
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                                                                                                                                                                                                                                                                                                                         Conservative
           GEIGY AG
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(first entry)
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93US-00139913
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                                                    94WO-IB000359
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "Tyr(2,3,5,6-F4)
                                                                                                                                                                                                                                                                                                                                   86.7%;
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Pred. No. 1.7e
0; Mismatches
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Best Local S
Matches 2
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10-OCT-2001;
18-DEC-2001;
The present invention describes a human leukocyte antigen (HLA) (e.g. soluble HLA) ligand database assembled by a process which involves providing a computer system capable of storing HLA data as a database, producing HLA having ligands loaded on it, isolating the loaded ligands from HLA, sequencing the loaded ligands to obtain HLA ligand data, and populating the database with HLA ligand data. Also described: (1) accessing HLA (e.g. sHLA) data stored in a database, and (2) a computer system for HLA (e.g. sHLA) ligand database. The database is populated with HLA ligand sequences, motifs, extended motifs, submotifs, ligands
                                                                                                                                             Human leukocyte antigen ligand database assembled by producing HIA loaded ligands, isolating and sequencing loaded ligands to obtain I ligand data and populating database in computer system with ligand
                                                                                                                                                                                                                                               (HILD/)
(PRIL/)
(HICK/)
                                                                                                                         Disclosure; Fig 5; 118pp; English.
                                                                                                                                                                                                                       Hildebrand WH, Prilliman
                                                                                                                                                                                                                                                                                                                                           21-FEB-2002; 2002WO-US005298.
                                                                                                                                                                                                 WPI; 2002-732755/79.
                                                                                                                                                                                                                                                                                                                                                                                           WO200269198-A2
                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Polypeptide receptor antagonists to tetrapeptide (AAR75498 or AAR75499)
                                                                                                                                                                                                                                                                                                                                                                                                                                         ligand;
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                                                                                                                                                                                                                                              HILDEBRAND W H. PRILLIMAN K R. HICKMAN H D.
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2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                         human
major
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; 2001US-00974366.
; 2001US-00022066.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sHLA molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Boyar
                                                                                                                                                                                                                                                                                                                                                                                                                                       leukocyte antigen; histocompatibility
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    86.7%;
66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antagonists to C5a contain
8 or AAR75499) or core trip
activity. (Updated on 25-N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ligand related peptide
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Pred. No. 1.7e+06;
O; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8
                                                                                                                                                                                                                       Hickman
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               agonist activity - are used 
eases and inflammatory condit
                                                                                                                                                                                                                                                                                                                                                                                                                                       predictive algorithm; database;
complex; viral; bacterial; tumour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NG,
                                                                                                                                                                                                                       HU;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         tain an essential core
tripeptide (DGA) which
25-MAR-2003 to correct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          #7.
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d data.
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RESULT 6
ABP55796
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Best Local S
Matches 2
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MHC ligand;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               unique to infected cells, tumour specific ligands, as well as a collection of current and future developed HIA ligand sequences. The database which contains endogenously bound and loaded ligands facilitates searching of viral, bacterial, tumour or human protein sequences for ligands likely to bind a particular HIA class I or class II protein. The database allows the user to screen an unknown peptide sequence for potential matches with sHIA ligand discrete sequences or sHIA ligand motifs of sequences. Due to the completeness and concentration of sHIA obtained to date, better sequencing data of numerous endogenously loaded HIA ligands is found in the sHIA ligand database, and by comparison of such ligands to each other and to the genomic sequence, better motifs are also found in the sHIA ligand database. ABPS592 to ABPS912 represent amino acid sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                    21-FEB-2001; 2001US-0270357P.
10-OCT-2001; 2001US-00974366.
18-DEC-2001; 2001US-00022066.
                                                                                                                                                                                                                            Hildebrand
                                                                                                                                                                                                                                                   (HILD/) HILDEBRAND W
(PRIL/) PRILLIMAN K
(HICK/) HICKMAN H D.
                                                                                                                                                                                                                                                                                                                                                      21-FEB-2002; 2002WO-US005298
                                                                                                                                                                                                                                                                                                                                                                                 06-SEP-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               B15 class I
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 4 AA;
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                                                                                                                                                                                                                            HW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                           human
major
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SHLA molecule ligand related peptide #90.
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                                                                                                                                                                                                                            Prilliman
                                                                                                                                                                                                                                                                                                                                                                                                                                                         leukocyte antigen; predictive algorithm; datal
histocompatibility complex; viral; bacterial;
                                                                                                                                                                                                                                                                X R H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    86.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Score 13; DB;
; Pred. No. 1.7e
0; Mismatches
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1.7e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
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The present invention describes a human leukocyte antigen (HLA) (e.g. soluble HLA) ligand database assembled by a process which involves providing a computer system capable of storing HLA data as a database, producing HLA having ligands loaded on it, isolating the loaded ligands from HLA, sequencing the loaded ligands to obtain HLA ligand data, and populating the database with HLA ligand data. Also described: (1) accessing HLA (e.g. sHLA) data stored in a database; and (2) a computer system for HLA (e.g. sHLA) ligand database. The database is populated with HLA ligand sequences, motifs, extended motifs, submotifs, ligands

Human leukocyte antigen ligand database assembled by producing HLA have loaded ligands, isolating and sequencing loaded ligands to obtain HLA ligand data and populating database in computer system with ligand database in computer system with ligand database.

data.

WPI; 2002-732755/79.

Disclosure; Fig 5; 118pp;

English.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Soluble human leukocyte antigen; HLA; sHLA; cell pharm; multimeric HLA complex; cytotoxic T lymphocytes; CTL; immune response; viral infection; cancer; autoimmune disease; vaccine development; MHC; major histocompatibility complex; diagnostic development; HLA-class I polymorphism; HLA-B15 allotype; ligand.
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The invention describes a method of producing soluble human leukocyte antisen (HIA) molecules (sHIA) in cell pharm involving amplifying HIA allelic DNA by PCR using a locus specific primer to produce truncated a PCR product (PI), inserting P1 into mammalian expression vector; electroporating the plasmid into a host cell; inoculating the cell pharm with the host cell such that cell pharm produces sHIA. A multimeric HIA complex (I) is useful for testing functionality of peptide ligands bound
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10-OCT-2001; 2001US-00974366.
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RESULT 8
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                                                                                        Producing soluble human leukocyte antigen (HLA) in cell pharm useful studies of peptide loading for characterizing human immune responses involves using HLA allelic cDNA or genomic DNA as starting material.
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Disclosure; Page 183;

300pp; English

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CC The invention describes a method of producing soluble human leukocyte CC antigen (HLA) molecules (sHLA) in cell pharm involving amplifying HLA CC allelic DNA by PCR using a locus specific primer to produce truncated a PCR product (PI), inserting P1 into mammalian expression vector; CC electroporating the plasmid into a host cell; inoculating the cell pharm coulet ship the cell such that cell pharm produces sHLA. A multimeric HLA CC complex (I) is useful for testing functionality of peptide ligands bound by at least two soluble HLA molecules. (I) can be tested for its ability to serve as ligands for cytotoxic Tlymphocytes (CTLs) and induce immune to serve as ligands for cytotoxic Tlymphocytes (CTLs) and induce immune couleting the human immune system to induce tolerance in autoimmune couleting the human immune system to induce tolerance in autoimmune couleting the human immune system to induce tolerance in autoimmune couleting the human immune system to induce tolerance in autoimmune couleting the human immune system to induce tolerance in autoimmune couleting the human immune system to induce tolerance in autoimmune couleting the human immune system to induce tolerance in autoimmune couleting the individual secreted major histocompatibility complex (MHC) can be produced the secreted MHC molecules are also useful to generate ligands and hence couleted class I HLA-BL5 allotypes; compare the different ligand maps to identify potentially shared elements; and characterise the elements coulet identified to positively or negatively validate the courrence of coverlapping ligands. The truncated version of (MHC) can be produced in mammalian or insect/bacterial cells such that milligram or greater couleting the course of the courrence of course couleting the course couleting that milligram or greater course couleting the course cours
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18-DEC-2000; 2000US-0256410P.
10-OCT-2001; 2001US-00974366.
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                                      Hildebrand WH,
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Sequence 4 AA;

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CC antigen (HLA) molecules (sHLA) in cell pharm involving amplifying HLA CC allelic NAA by PCR using a locus specific primer to produce truncated a CC PCR product (PI), inserting P1 into mammalian expression vector; CC electroporating the plasmid into a host cell; inoculating the cell pharm (CC with the host cell such that cell pharm produces sHLA. A multimeric HLA CC complex (I) is useful for testing functionality of peptide ligands bound CC by at least two soluble HLA molecules. (I) can be tested for its ability CC to serve as ligands for cytotoxic T lymphocytes (CTLs) and induce immune CC gathological conditions such as viral infections and cancer, and for CC modulating the human immune system to induce tolerance in autoimmune CC diseases. The individual secreted major histocompatibility complex (MHC) CC molecules produced are useful for studies of peptide loading (i.e., in CC vaccine development) and to the development of diagnostics. With the CC secreted MHC molecules, naturally loaded peptides can be eluted from the sessement of structural and functional impact of HLA class I polymorphism. The molecules are also useful to generate ligands and hence CC ligand maps from the peptide pools extracted from series of distinct yet crelated class I HLA-BL5 allotypes; compare the different ligand maps to CC identified to positively or negatively validate the occurrence of coverlapping ligands. The truncated version of (MHC) can be produced in CC quantities of an individual class I or class I molecule can be obtained. CC flass sequence expresents a HLA (human language article ligand negatively validate the occurrence of operator article ligand.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention antiqen (HLA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Producing soluble human leukocyte antigen (HLA) in cell pharm useful studies of peptide loading for characterizing human immune responses involves using HLA allelic cDNA or genomic DNA as starting material.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Fig 26; 300pp; English.
                                                      sequence
                                                   represents a HLA (human leukocyte antigen) peptide ligand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      describes a method of producing soluble human molecules (sHLA) in cell pharm involving ampli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Leukocyte
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RESULT 10
ABP67652
ID ABP67
XX ABP67
AC ABP67
AC ABP67
XX IO-DE
XX Human
XX Human
XX Immun
XX Immun
XX IO-DE
XX Antib
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XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; CD66; CEACAM; cytostatic; antiinflammatory; immunomodulator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABP67652 standard; peptide; 4 AA
                                                                                                                                                                                                                                                                                                                                                                             WO200268601-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immune system;
inflammatory di
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antibacterial; virucide; gene therapy; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-DEC-2002
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2; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         disease;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          autoimmune
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 bacterial; virus;
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28-FEB-2001; 2001US-0272113P 27-FEB-2002; 2002WC-US005720

(SKUB/) SKUBITZ

AM.

06-SEP-2002

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RESULT 11
ABP67653
ID ABP677673
XX ABP677
XX ID-DE
XX Humar
XX Humar
XX Humar
XX Humar
XX Humar
XX Homo
XX Homo
XX Infla
XX 27-FI
XX 28-FI
XX 28-FI
XX X ESKUI
PA (SKUI
PA (SK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to an isolated peptide (I) from a surface exposed cregion of a CD66 family member (ABB671307). The peptide CC modulates: (a) activation of neutrophils; (b) activation or inhibition of CC modulates: (a) activation of neutrophils; (b) activation or inhibition of T-cells, B-cells, Mc cells, LAK cells, dendriic cells or other immune CC system cells; (c) proliferation and/or differentiation of the above CC cells, including epithelial cells; (d) homotypic and/or heterotypic activation and/or members; and (e) adhesion of CD66 family members; and (e) adhesion of CD66 family the CC function of CD66 family members and/or their ligands, such as activation CC function of CD66 family members and/or their ligands, such as activation CC fufferentiation of the immune cells. The peptides may also be used in treating or diagnosing autoimmune diseases, cancer, infections (e.g. C therapies and for immunisation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; CD66; CEACAM; cytostatic; antiinflammatory; immunomodulator; antibacterial; virucide; gene therapy; vaccine; neutrophil; immune system; autoimmune disease; cancer; infection; bacterial; vi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 4 AA;
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New peptide from a surface exposed region of a CD66 family member, useful for modulating the function of CD66 family members, e.g. activation of neutrophils, for treating or diagnosing autoimmune diseases or cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  neutrophils,
                                                                                                                             WPI; 2002-706981/76
                                                                                                                                                                                                                                                                                                                                                                                                 27-FEB-2002; 2002WO-US005720.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABP67653 standard;
                                                                                                                                                                                                                                                                                                                                    28-FEB-2001; 2001US-0272113P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200268601-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        inflammatory disease; transplantation; immunisation
                                                                                                                                                                                                                                                                          (SKUB/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CD66 family modulating peptide SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        a vaccine.
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3
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SKUBITZ A P N.
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2; Conser
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                                                                                                                                                                                      Skubitz APN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   86.7%;
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Pred. No. 1.7e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 bacterial; virus;
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AAU84
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AC Caspa
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                                                                                                     Novel apoptosis-associated and/or -modified protein useful as the diagnosis, prevention or treatment of apoptosis-associated particularly neurodegenerative diseases.
                                                                                                                                                                                                                                                                                     16-JUN-2000; 2000EP-00112813.
16-NOV-2000; 2000EP-00125013.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Caspase cleavage site #11.
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                                                                    Claim 30; Page 62; 86pp; English.
                                                                                                                                                                               WPI; 2002-098088/13
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                                                                                                                                                                                                                                                    (PLAC) MAX PLANCK GES FOERDERUNG WISSENSCHAFTEN
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2; Conserv
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                                                                                                                                                                                                                8
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66.7%;
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Pred. No. 1.7e
0; Mismatches
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1.7e+06;
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                                                                                                                                as target for
ated diseases
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The invention relates to an apoptosis-associated and/or -modified protein (I) selected from GAP SH3 binding protein, HCD2 and AOP-1 (not defined) or its proteolytic fragments. (I) is useful as target for the diagnosis,

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Gaps

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RESULT 13
ABJ18638
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Best Local &
Matches 2
                                                                                                                                       New peptide antagonists against etreating tumors or cancers (e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  prevention or treatment of apoptosis-associated diseases, and in a method for identifying apoptosis modulators. (I) is useful for treatment of hyperproliferative or degenerative diseases including cancers, autoimmune diseases, Alzheimer's disease, viral infections such as acquired infarction. (I) is also useful for treating and/or preventing diseases related to oxidative stress like ischaemic stroke, arthritis, heart failure, Parkinson's disease and amyotrophic lateral sclerosis. (I) is also useful for develop modification-specific diagnostic tools. (I) is also useful for developing protein chips or other solid phase screening devices for high throughput screens, and also as a diagnostic tool to screen for caspase activity, and/or to determine the effectivity of caspase cleavage inhibiting and/or modulating substances. AAU84237-AAU84260 represent caspase cleavage sites of the invention
                                                                                                                                                                               WPI; 2003-075482/07
                                                                                                                                                                                                                     Greene MI,
                                                                                                                                                                                                                                                                       06-APR-2001;
03-AUG-2001;
                                                                                                                                                                                                                                                                                                              08-APR-2002; 2002WO-US011211.
                                                                                                                                                                                                                                                                                                                                                                                              Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                               Vaccine; erbB receptor antagonist; tumour; cancer; gastric cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           ErbB receptor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    prevention or treatment of apoptosis-associated diseases, for the manufacture of a pharmaceutical agent and for identifying apoptos modulators. The protecome is useful as a target for the diagnosis,
                                                                                                                                                                                                                                                                                                                                                                   WO200281649-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABJ18638 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                      cancer;
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2; Conser
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2001US-0309864P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          antagonist peptide #21
                                                                                                                                                                                                                                                                                                                                                                                                                      prostate cancer
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Pred. No. 1.7e+06;
0; Mismatches 1
                                                                                                                                       erbB receptors, useful lung adenocarcinomas,
                                                                                                                                                                                                                     Richter
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                                                                                                                                                                                                                     Berezov
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The invention comprises peptide antagonists designed to target erbB receptors. The erbB receptor peptide antagonists are useful for preventing tumours and cancers (e.g. gastric cancer, breast cancer and prostate cancer). The present amino acid sequence represents an erbB receptor peptide antagonist of the invention

RESULT 15 ADJ81984

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ADJ81984;

ADJ81984 standard; peptide; 4 AA.

Claim 1; or prostate

Page

82; 115pp; English.

cancer)

in humans

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for preventing or breast carcinomas
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                                                                                                                                                                                                                                                         The present invention relates to novel antibodies, which have specifor activated cell surface receptors (erbB, TNF and immunoglobin ge superfamily: 19SF) and bind to an epitope (ABP59223-ABP59259). The antibodies are useful for preventing transformation of a normal cel a tumour cell in a patient who had a tumour with tumour cells that express activated erbB receptors on the cell surface surgically rem or in whom the tumour has entered remission, for treating a patient suffering from a cancer characterised by tumour cells that their cell surfaces, or for treating a patient suffering from an er tumour. The antibodies were produced by immunising a suitable host peptide (ABP59223-ABP59259), or its reverse sequence
                                                                                                                                                                                                                   Sequence 4 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel antibody for preventing, treating, imaging or diagnosing tumor, I specificity for activated cell surface receptor and binds specifically an epitope that does not comprise phosphotyrosine residue.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 53; Page 88; 114pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-268117/26
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immunoglobin
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2; Conser
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                                                                                                      Conservative
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                                                                                                                                 Score 13; DB 6;
Pred. No. 1.7e+06;
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0; Mismatches
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IgSF; epitope.
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Search completed: December 13, 2004, 22:22:40 Job time : 80 secs
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                                                                                                    Query Match
Best Local Similarity 60.,
Thehes 2; Conservative
                                                                                                                                                                                                                      The invention relates to novel nucleic acid polymerases from the thermophilic organism Thermus brookianus or mutants having a mutation that decreases 5'-3' exonuclease activity or reduces discrimination against dideoxymucleotide triphosphates. The nucleic acid polymerases, isolated nucleic acids encoding the polymerases, vectors, host cells and kits are useful for DNA synthesis, primer extension, DNA sequencing, reverse transcription, or DNA and RNA amplification procedures. The nucleic acids are also useful for the recombinant expression of the polymerase polymerase polypeptides. The nucleic acid polymerases and nucleic acids are useful for the detection and isolation of nucleic acids encoding DNA polymerase I activity. This sequence corresponds to a peptide of a polymerase of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New Thermus brockianus nucleic acid polymerases, useful for DNA synthesis, primer extension, DNA sequencing, reverse transcription, or DNA and RNA amplification procedures.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Thermus brockianus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acid polymerase; thermophile; mutation; exonuclease activity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-JUN-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Thermus brockianus nucleic acid polymerase peptide fragment
                                                                                                                                                                                             Sequence 4 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rozzelle J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-NOV-2001; 2001US-0334434P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-NOV-2002; 2002WO-US037657.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; SEQ ID NO 18; 74pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-505285/47.
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Match
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2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

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Copyright (c) 1993 - 2004 Compugen Ltd.
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            US-08-798-897-49
US-08-463-224-2
US-08-463-377-2
US-08-978-523-49
US-08-978-523-49
US-09-978-523-49
US-09-978-523-49
US-09-978-523-49
US-09-978-528-36
US-08-628-68-36
US-08-628-68-66-14
US-08-618-696-14
US-08-618-696-14
US-08-67-974-36
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US-08-467-976-36
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Sequence 49, Appli Sequence 2, Appli Sequence 2, Appli Sequence 199, Appl Sequence 199, Appl Sequence 199, Appl Sequence 199, Appl Sequence 36, Appli Sequence 36, Appli Sequence 30, Appli Sequence 31, Appli Sequence 36, Appli Sequence 37, Appli Sequence 37, Appli Sequence 38, Appli Sequence 39, Appli Sequence 31, Appli Sequence 21, Ap
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; MOLECULE TYPE: peptide US-08-798-897-49

Query Match
Best Local Similarity
Matches 2; Conserv

Conservative

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Score 13; DB 1; Pred. No. 3.8e+05; 0; Mismatches 1

Length 4; Indels

0;

Gaps

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86.7%;

US-08-798-897-49 (Sequence 49, Application US Patent No. 5789201 (GENERAL IMPORNATION: General INFORNATION: GUSTITLE OF INVENTION: General INFORMATION: General INFORMATION: General INFORMATION: General INFORMATION: General INFORMATION: How York CITY: Washington STATE: DC COUNTRY: USA ZIP: 2005 COMPUTER RADABLE FORM: MEDIUM TYPE: Ploppy d COMPUTER RADABLE FORM: MEDIUM TYPE: Ploppy d COMPUTER PAPLICATION UNUBER: CURRENT APPLICATION UNUBER: GENERAL APPLICATION UNUBER: GENERAL INFORMATION INFORMATION INFORMATION INFORMATION OF SEQUENCE CHARACTERISTICS INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS LENGTH: 4 maino acids STRANDEDNES: not rel TOPOLOGY: linear	28 13 86.7 29 13 86.7 30 13 86.7 31 13 86.7 32 13 86.7 34 13 86.7 35 13 86.7 36 13 86.7 37 13 86.7 38 13 86.7 40 13 86.7 41 13 86.7 42 13 86.7 43 13 86.7 44 13 86.7 45 13 86.7 46.7 47 13 86.7 48 13 86.7 49 13 86.7 40 13 86.7 41 13 86.7 42 13 86.7 43 13 86.7 44 13 86.7
on US/08798897 la, John Genes Coding For Ecl-y, a Ecl- Homologue : 53 ESS: E. KESSLER, GOLDSTEIN & FOX P.L. York Avenue, N.W., Suite 600 PORTA: Compatible CC-DOS/MS-DOS IN Release #1.0, Version #1.30 DATA: US/08/798,897 ruary 11, 1997 435 RRANION: bert W. BR: 32,893 NUMBER: 1483.0140001 NORMATION: 11-2600 -2540 DNO: 49: BCIGS: STICS: STICS: STICS: STICS: SCICS: STICS: CRESTICS: CRE	6 1 US-08-248-538-40 6 1 US-08-260-199A-31 6 1 US-08-260-199A-32 6 1 US-08-260-199A-32 6 1 US-08-269-050-21 05-08-461-955-21 05-08-461-955-21 05-08-978-523-53 0 US-08-978-523-53 0 US-08-932-787B-12 0 US-08-932-787B-12 0 US-08-932-012C-12 0 US-09-518-737-6 0 5200320-19 7 1 US-09-541-997A-19 7 2 US-09-070-756-8 7 3 US-09-193-104-28
e e e	Sequence 40, Appl Sequence 31, Appl Sequence 32, Appl Sequence 53, Appl Sequence 14, Appl Sequence 21, Appl Sequence 21, Appl Sequence 12, Appl Sequence 13, Appl Sequence 19, Appl Sequence 19, Appl Sequence 19, Appl Sequence 28, Appli Sequence 28, Appli

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US-08-463-377-2
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Matches 2; Conserv
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                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 4 amino acids
                                                                                                                                                                                                                                       APPLICANT: van Oo:
APPLICANT: Boyar,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: FOLEY, Shawn P.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 908-654-5000
TELEPAX: 908-654-7866
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: van Heeke, Gino
TITLE OF INVENTION: C5a Receptor Antagonists Having
TITLE OF INVENTION: Substantially No. 5807824Agonist Activity
NUMBER OF SEQUENCES: 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                               STREET: 600 --
STREET: Westfield
CITY: Westfield
CTATE: NJ
TCA
                                                                                NUMBER OF SEQUENCES: 67
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lerner, David, Littenberg, Krumholz & Mentlik
STREET: 600 South Avenue West
                                                                                                                                      APPLICANT: van Heeke, Gino
TITLE OF INVENTION: C5a Receptor Antagonists Having
TITLE OF INVENTION: Substantially No. 5837499Agonist Activity
NUMBER OF SEQUENCES: 67
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 50.
TELEFAX: 139-125
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ZIP: 07090
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE:
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                                                                                                                                                                                                                                                                                                           Application US/08463377
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600 South Avenue West
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      USA
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Schmitz, Albert
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Galakatos, Nicholas G.
Schmitz, Albert
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                                                                                                                                                                                                                                                             van Oostrum, Jan
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Pred. No. 3.8e+05;
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US-08-978-523-49
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Best Local Similarity oo...
""" ches 2; Conservative
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                                                                                                       CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION DATA: US 08/798,897
APPLICATION UNMBER: US 08/798,897
FILING DATE: February 11, 1997
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Esmond, Robert W.
NAME: Esmond, Robert W.
AEGISTRATION NUMBER: 32,893
REGISTRATION NUMBER: 32,893
REGISTRATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                  TELEFAX: 202-371-2540 INFORMATION FOR SEQ ID NO:
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Foley, Shawn P.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 908-654-5000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IRM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/463,377
                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: TITLE OF INVENTION: NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                  SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX:
TYPE: amino acid
STRANDEDNESS: no
                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/978,523 FILING DATE: herewith
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Washington
STATE: DC
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                                 ENGTH:
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GY: linear
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                                   4 amino acids
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not relevant
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Pred. No. 3.8e+05
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                                                        Query Match
Best Local S
Matches 2
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                                                                                                                                                                                     NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 4 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN DATA:
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 35,238
REFERENCE/DOCKET NUMBER: 00
TELECOMMUNICATION INFORMATION:
TELECPHONE; (617) 542-5070
                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 08/227,360 FILING DATE: April 13, 1994 ATTORNEY/AGENT INFORMATION:
                                                                                                                                MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 536
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Yu, Guo-Liang
TITLE OF INVENTION: RPS2 GENE FAMILY, PRIMERS, PROBES, AND DETECTION
TITLE OF INVENTION: METHODS
                                                                                                                                              TOPOLOGY:
                                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                        TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/310,9
FILING DATE: September 22, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
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2 GTY 4
                                                      Similarity 66. 2; Conservative
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225 Franklin Street
                                                                                                                                                                                                                                               (617) 542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
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Brent, Andrew F.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mindrinos, Michael N.
                                                                                                                                                                                                                                                                                              | INFC:
| Karen F. | 35,238 | 007
                                                                                                                             protein
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                                                                                                                                                          not relevant
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                                                                   86.7%;
66.7%;
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                                                     Score 13; DB
Pred. No. 3.8e
0; Mismatches
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Pred. No. 3.8e+05
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                                                                 3.8e+05;
                                                                                  DB 2;
                                                                               Length 4;
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; ORGANISM: Arabidopsis thaliana
US-09-301-085-199
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GENERAL INFORMATION:
APPLICANT: Trowbri
APPLICANT: Collawn
APPLICANT: Tainer,
APPLICANT: Kuhn, L
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Best Local
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Patent No. 6262248
GENERAL INFORMATION:
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CURRENT FILING DATE: 1999-04-28
EARLIER APPLICATION NUMBER: 08/310,912
EARLIER FILING DATE: 1994-09-22
EARLIER APPLICATION NUMBER: 08/227,360
EARLIER FILING DATE: 1994-04-13
NUMBER OF SEQ ID NOS: 208
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ·US93-01669-41
                                                                                    PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US/07/844,852
                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: PCT/US93/01669
FILING DATE: 01-MAR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TILE OF INVENTION: RPS2 GENE FAMILY, PRIMERS, PROBES, AND ITLE OF INVENTION: DETECTION METHODS
                                                        APPLICATION NUMBER: US/0
FILING DATE: 03-MAR-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                         STREET: 1000
CITY: Los Angeles
CTATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Spensley Horn Jubas & Lubitz
STREET: 1880 Century Park East, Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                             UMBER OF SEQUENCES:
NAME: Wetherell, Jr. Ph.D., John R. REGISTRATION NUMBER: 31,678
REFERENCE/DOCKET NUMBER: PD-1636
                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TLE OF INVENTION:
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2; Conserv
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Collawn, ur.,
Tainer, John A.
Tainer, Leslie A.
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Pred. No. 3.
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PCT-US95-04589-199
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INFORMATION FOR SEQ ID NO: 41:
SEQUENCE CHARACTERISTICS:
LENGTH: 4 amino acid
                                                                         PCT-US95-04589-199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 66.7
Matches 2; Conservative
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         Matches
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APPLICATION NUMBER: US 08/227,360
APPLICATION NUMBER: US 08/227,360
APPLICATION NUMBER: 1994
ATTORNEY/AGENT INFORMATION:

NAME: Clark, Paul T.

REGISTRATION NUMBER: 30,162
REGISTRATION NUMBER: 00786/230001
TELECOMMUNICATION INFORMATION:
TELEPAN: (617) 542-5070
TELEPAN: (617) 542-8906
                                      Query Match
                                                                                                                                                        INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 4 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 02110-2904
COMPUTER FRADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30B
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Yu, Guo-Liang
TITLE OF INVENTION: RPS2 GENE AND USES THEREOF
NUMBER OF SEQUENCES: 201
CORRESPONDENCE ADDRESS:
                                                                                         TOPOLOGY: 1i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Ausubel, Frederick M. APPLICANT: Staskawicz, Brian J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 225 F
       Local Similarity hes 2; Conserv
                                                                                                          STRANDEDNESS: not TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE: MA
                                                                                                                                               TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
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225 Franklin Street Suite 3100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application PC/TUS9504589
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Dahlbeck, Douglas
Katagiri, Fumiaki
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Mindrinos, Michael N.
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       Conservative
                                                                                        protein
                                                                                                                            not relevant
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                         86.7%;
66.7%;
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Pred. No. 3.8e+05;
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         0; Mismatches
                         Score 13; DB 5;
Pred. No. 3.8e+05;
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                                          Length 4;
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US-07-962-569A-6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TOPOLOGY: linear; MOLECULE TYPE: protein US-07-962-569A-6
                                                                                                                                                                                                                                                         RESULT 10
US-08-292-968-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
ZIP: 92127
ZIP: 92127
COMPUTER READABLE FORM:
WEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
SOFTWARE: DatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/962,569A
FILING DATE: 19921013
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 31,286
REFERENCE/DOCKET NUMBER: 920224.01
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                      Sequence 30 Patent No.
                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local (
                                                                     APPLICANT: READ, Ranuy U.
APPLICANT: STEIN, Penelope E.
APPLICANT: COCKLE, Stephen A.
APPLICANT: COCKEN, Raymond P.
APPLICANT: LOOSMORE, Sheena
APPLICANT: KLEIN, Michel H.
APPLICANT: ARMSTRONG, Glen D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (619) 451-1120
TELEFAX: (619) 451-9628
INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: CHANG, YIN
APPLICANT: HAM, RICHA
TITLE OF INVENTION: H
NUMBER OF SEQUENCES:
         APPLICANT: ARMSTRONG, Glen D.
APPLICANT: HAZES, BART
TITLE OF INVENTION: MODIFICATION OF PERTUSSIS TOXIN
NUMBER OF SEQUENCES: 46
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: JEFFERS, KATHLEEN F. APPLICANT: CHANG, YING-FON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: MENON,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 11545 W. CITY: SAN DIEGO STATE: CA
ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
                                                                                                                                                                                                                          30, Application US/08292968
5. 5856122
                                                                                                                                                                                                                                                                                                                                                                                            Similarity 66.
2; Conservative
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11545 W. BERNARDO COURT, STE. 302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HAM, RICHARD G.
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Pred. No. 3.8e+05;
0; Mismatches 1
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Gaps

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RESULT 11
US-08-292-968-36
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APPLICANT: STEIN, Penelope E.
APPLICANT: COCKLE, Stephen A.
APPLICANT: COCKLE, Raymond P.
APPLICANT: LOOSMORE, Sheena
APPLICANT: LOOSMORE, Sheena
APPLICANT: KLEIN, Michel H.
APPLICANT: ARMSTRONG, Glen D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/110,947
FILING DATE: 24-AUG-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/251,121
FILING DATE: 31-MAY-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: STEWART, Michael I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-388
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1153
TELEPAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 30:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/292,968
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MEDIUM TYPE: Floppy disk
COMPUTER: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/292,968
                                                                                                                                                                                                                                 TITLE OF INVENTION: MODIFICATION OF PERTUSSIS TOXIN NUMBER OF SEQUENCES: 46
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
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STRANDEDNESS: sin
                                                                                                                                                                   STATE:
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                                                                                                                                             COUNTRY:
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M5G 1R7
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                                                                                                                                                                              Toronto
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                                                                                                                                                                                                E: Sim & McBurney
Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                         HAZES, Bart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 13; DB 2;
Pred. No. 3.8e+05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1; Indels
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RESULT 12
US-08-618-696-3
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TOPOLOGY:
US-08-292-968-36
                                  NAME: PARKER, DAVID L.

REGISTRATION NUMBER: 32.165

REGISTRATION NUMBER: 32.165

REPERENCE DOCKET NUMBER: UOAB:002/PAR

TELECOMMUNICATION INFORMATION:

TELEPHONE: 512-320-7200

TELEPAX: 512-474-7577

TELEPX: NOT APPLICABLE
INFORMATION FOR SEQ ID NO: 3:

SEQUENCE CHARACTERISTICS:

SEQUENCE CHARACTERISTICS:

TENER: 5 amino acid residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 3, Application Patent No. 5861475
GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/618,696
FILING DATE: 20-MAR-1996
CLASSIFICATION: 514
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PRIOR APPLICATION DATA:
APPLICATION UMMBER: US 08/251
PILING DATE: 31-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Michael I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (416) 595-116
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: US 08/110,947
APPLICATION NUMBER: US 08/110,947
                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/995,269
FILING DATE: 12/21/9
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: COOPER, Jr., J. ALLEN D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: INHIBITION OF PHAGOCYTES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD, WH.
STREET: P.O. BOX 4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: P.O. B
CITY: HOUSTON
STATE: TEXAS
LENGTH: 5 amino acid residues TYPE: amino acid stranDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: U
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20-MAR-1996
V. 517
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Pred. No. 3.8e+05;
0; Mismatches 1
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RESULT 14
US-08-467-974-30
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TOPOLOGY:
US-08-618-696-3
                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 514

PRIOR APPLICATION DATN:
APPLICATION UNMBER: 07/995,269

FILING DATE: 12/21/92

ATTORNEY/AGENT INFORMATION:
NAME: PARKER, DAVID L.
REGISTRATION NUMBER: 32,165
REFERENCE/DOCKET NUMBER: UOAB:002/PAR
TELECOMMUNICATION: 1070RMATION:
TELEPHONE: 512-320-720
TELEPHONE: 512-474-7577
TELEEX: NOT APPLICABLE
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
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Sequence 30, Application US/08467974
Patent No. 5965385
GENERAL INFORMATION:
APPLICANT: READ, Randy J.
APPLICANT: STEIN, Penelope E.
APPLICANT: COCKLE, Stephen A.
                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 14, Application US/08618696 Patent No. $861475 GENERAL INFORMATION: APPLICANT: COOPER, Jr., J. ALLEN
                                                                                                                                                                                                                                                                  Best Local Similarity
                                                                                                                                                                                                                                                                                     Query Match
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORDPERPECT 5.1
CURENT APPLICATION DATA:
APPLICATION NUMBER: US/08/618,69
FILING DATE: 20-MAR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: COOPER, Jr., J. ALLEN D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS
TITLE OF INVENTION: INHIBITION OF PHAGOCYTES
NUMBER OF SEQUENCES: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: FLOPPY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: US/
                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 66.7 hes 2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: ARNOLD, WH
STREET: P.O. BOX 4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: HOUSTON
                                                                                                                                                                                                                                                                                                                                                                                            ENGIH:
                                                                                                                                                                                                                  1 GXY 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 GXY 3
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                                                                                                                                                                                                                                                                                                                                                                                            5 amino acid residues
                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FLOPPY DISK
                                                                                                                                                                                                                                                                                                                                                        single
                                                                                                                                                                                                                                                                    86.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   86.7%;
66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WHITE & DURKER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US/08/618,696
                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 13; DB 2;
Pred. No. 3.8e+05;
                                                                                                                                                                                                                                                                    Score 13; DB 2;
Pred. No. 3.8e+05;
                                                                                                                                                                                                                                                    1; Indels
                                                                                                                                                                                                                                                                                     Length 5;
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US-08-467-974-36
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                                                                                      Patent No. 5965385
GENERAL INFORMATION:
APPLICANT: READ, RA
APPLICANT: STEIN, I
APPLICANT: COCKLE,
APPLICANT: LOOSMORE
APPLICANT: LOOSMORE
                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-467-974-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467 and PATENTAL DATE:
                                                                                                                                                                                                                Sequence 36,
                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/292,968
FILING DATE: 22-AUG-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/251,121
FILING DATE: 31-MAY-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/110,947
APPLICATION DATA:
24 ANG-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (416) 595-11
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: STEWART, Michael I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-454 MIS:vg
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
TELEPAX: (416) 595-1163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/467,536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 24-AUG-1993
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Michael I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: Canada
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: HAZES, Bart
TITLE OF INVENTION: MODIFICATION OF PERTUSSIS TOXIN
NUMBER OF SEQUENCES: 46
                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 5 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: Suite
CITY: Toronto
                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
   OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                         1 GXY 3
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                                                                                                                                                                                                                Application US/08467974
                                                 STEIN, Penelope E. COCKLE, Stephen A. OOMEN, Raymond P. LOOSMORE, Sheena KLEIN, Michel H. ARMSTRONG, Glen D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: Sim & McBurney
Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                      HAZES, Bart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  KLEIN, Michel H.
ARMSTRONG, Glen D.
                                                                                                                                                              READ, Randy J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OOMEN, Raymond P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOOSMORE, Sheena
                                                                                                                                                                                                                                                                                                                                                                                            86.7%;
66.7%;
   MODIFICATION OF PERTUSSIS TOXIN 46
                                                                                                                                                                                                                                                                                                                                                                         Score 13; DB 2;
Pred. No. 3.8e+05;
0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                             DB 2; Length 5;
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                                                                                                                                                                                                                                                                                                                                                                             Gaps
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
CITY: Toronto
STATE: Ontario
COUNTRY: Canada
ZIP: McG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Lam PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
APPLICATION NUMBER: US/08/467,974
FILING DATE: O-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/467,536
FILING DATE: O-JUN-1995
APPLICATION NUMBER: US/08/292,968
FILING DATE: O-JUN-1994
PRIOR APPLICATION NUMBER: US/08/292,968
FILING DATE: 22-AUG-1994
PRIOR APPLICATION NUMBER: US/08/291,121
FILING DATE: 31-MAY-1994
PRIOR APPLICATION NUMBER: US/08/251,121
FILING DATE: 24-AUG-1993
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Michael I.
REGISTRATION NUMBER: US/08/454 MIS:vg
TELEPAN: (416) 595-1163
INFORMATION FOR SEQ ID NO: 36:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acidd
STRANDEDNESS: mino acidd
STRANDEDNESS: single
US-08-467-974-36
Search completed: December 13, 2004, 22:34:51 Job time: 21.4 secs
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                                                                                                                                                          Query Match 86.7%;
Best Local Similarity 66.7%;
Matches 2; Conservative
                                                                                                                          1 GXY 3
                                                                                  2 GTY 4
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Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BLOSUM62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-10-068-905-8
/15
1 GXYX 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         December 13, 2004, 22:31:47; Search time 68.6 Seconds (without alignments) 20.827 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1585576 segs, 357178320 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gapop 10.0 , Gapext 0.5
                                                                           11554:
                                                                                                                                                                                                                                                                                     GenCore version 5.1.6 Copyright (c) 1993 - 2004 Compugen Ltd.
/cgn2_6/prodata/2/pubpaa/US08_PUBCOMB.pep:*
/cgn2_6/prodata/2/pubpaa/US08_PUBCOMB.pep:*
/cgn2_6/prodata/2/pubpaa/US09A_PUBCOMB.pep:*
/cgn2_6/prodata/2/pubpaa/US09_PUBCOMB.pep:*
/cgn2_6/prodata/2/pubpaa/US09_PUBCOMB.pep:*
/cgn2_6/prodata/2/pubpaa/US10A_PUBCOMB.pep:*
/cgn2_6/prodata/2/pubpaa/US10B_PUBCOMB.pep:*
/cgn2_6/prodata/2/pubpaa/US10B_PUBCOMB.pep:*
/cgn2_6/prodata/2/pubpaa/US10_PUBCOMB.pep:*
/cgn2_6/prodata/2/pubpaa/US10_NEW_PUB.pep:*
/cgn2_6/prodata/2/pubpaa/US10_NEW_PUB.pep:*
/cgn2_6/prodata/2/pubpaa/US10_NEW_PUB.pep:*
/cgn2_6/prodata/2/pubpaa/US10_NEW_PUB.pep:*
/cgn2_6/prodata/2/pubpaa/US10_NEW_PUB.pep:*
/cgn2_6/prodata/2/pubpaa/US10_NEW_PUB.pep:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

13	12	11	10	ø	80	7	თ	ហ	4.	ω	N		Result No.
13	13	13	13	13	13	13	13	13	13	13	13	13	Score
86.7	86.7	86.7	86.7	86.7	86.7	86.7	86.7	86.7	86.7	86.7	86.7	86.7	Query
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9	9	9	16	16	16	14	14	10	16	15	14	9	B
US+09-990-762-41	US-09-990-762-40	US-09-990-762-36	US-10-394-471B-11	US-10-394-471B-3	US-10-738-809-8	US-10-376-121A-143	US-10-344-779-8	US-09-792-686A-34	US-10-613-765-199	US-10-613-472-199	US-10-302-817A-18	US-09-867-852-199	H D
Sequence 41, Appl	40,	Sequence 36, Appl	Sequence 11, Appl	Sequence 3, Appli	Sequence 8, Appli	Sequence 143, App	Sequence 8, Appli	Sequence 34, Appl	Sequence 199, App	Sequence 199, App	Sequence 18, Appl	Sequence 199, App	Description

ALIGNMENTS

```
APPLICANT: YU, GNO-ILANG
ITTLE OF INVENTION: RPS2 GENE FAMILY, PRIMERS, PROBES, AND
ITTLE OF INVENTION: DETECTION METHODS

PILE REFERENCE: 00786/254002
CURRENT APPLICATION NUMBER: US/09/867,852
CURRENT FILING DATE: 2001-05-29
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/301,085
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/301,085
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 08/310,912
PRIOR PILING DATE: BARLIER FILING DATE: 1994-09-22
PRIOR PILING DATE: EARLIER FILING DATE: 1994-04-28
PRIOR PILING DATE: EARLIER FILING DATE: 1994-04-13
NUMBER OF SEQ ID NOS: 208
PRIOR FILING DATE: SARLIER FILING DATE: 1994-04-13
                                                                                                                              ; ORGANISM: Arabidopsis thaliana US-09-867-852-199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 1
US-09-867-852-199
                                                                                                                                                                                                                                 SOFTWARE: FASTSEQ for Windows Version 4.0 SEQ ID NO 199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 199, Application US/09867852
Patent No. US20020147324A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
Query Match 86.7%; Score 13; DB 9; Length 4; Best Local Similarity 66.7%; Pred. No. 1.4e+06; Matches 2; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Ausubel, Frederick M. APPLICANT: Staskawicz, Brian J. APPLICANT: Brent, Andrew F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Katagiri, Fumiaki
Kunkel, Barbara N.
Mindrinos, Michael N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dahlbeck, Douglas
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1; Indels

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; ORGANISM: Arabidopsis thaliana US-10-613-472-199
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US-10-302-817A-18
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                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/613,472
CURRENT FILING DATE: 2003-07-02
PRIOR APPLICATION NUMBER: US 09/867,852
PRIOR FILING DATE: 2001-05-29
PRIOR FILING DATE: 2001-05-29
PRIOR FILING DATE: 10999-04-28
PRIOR PRIOR APPLICATION NUMBER: US 09/301,085
PRIOR APPLICATION NUMBER: US 08/310,912
PRIOR APPLICATION NUMBER: US 08/310,912
PRIOR APPLICATION NUMBER: US 08/227,360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18, Application US/10302817A

Publication No. US20030198978A1

GENERAL INFORMATION:
APPLICANT: ROZZELLE, James
APPLICANT: BOLCHAKOVA, Elena
TITLE OF INVENTION: THERMUS BROCKIANUS NUCLEIC ACID POLYMERASES
FILE REFERENCE 4768US
FULE REFERENCE 4768US
CURRENT APPLICATION NUMBER: US/10/302,817A
CURRENT PILING DATE: 2002-11-22
PRIOR APPLICATION NUMBER: 60/334,434
PRIOR FILING DATE: 2001-11-30
NUMBER OF SEQ ID NOS: 58
OURSENTANT SECONDAME.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Brent, Andrew F.
APPLICANT: Dahlbeck, Douglas
APPLICANT: Kantagiri, Funiaki
APPLICANT: Kunkel, Barbara N.
APPLICANT: Kunkel, Barbara N.
APPLICANT: Mindrinos, Michael N.
APPLICANT: YU, Guo-Liang
TITLE OF INVENTION: PS2 GENE FAMILY, PRII
TITLE OF INVENTION: DETECTION METHODS
FILE REFERENCE: 00786/254004
                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 214
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 199, Application No. US200 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn version 3.2 SEQ ID NO 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Ausubel, Frederick M.
APPLICANT: Staskawicz, Brian J.
APPLICANT: Brent, Andrew F.
APPLICANT: Dahlbeck, Douglas
APPLICANT: Katagiri, Pumiaki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Thermus brockianus
                                                                                                                                                                                       LENGTH: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            y Match
Local Similarity 66.7%;
  Local Similarity 66.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 GXY 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Application US/10613472
o. US20040088756A1
86.7%; Score 13; DB 15; Length 4; 66.7%; Pred. No. 1.4e+06; Live 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 13; DB 14; Length 4; Pred. No. 1.4e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIMERS,
     0
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Gaps
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CURRENT APPLICATION NUMBER: US/10/613,765
CURRENT FILING DATE: 2003-07-02
PRICR APPLICATION NUMBER: US 09/867,852
PRICR FILING DATE: 2001-05-29
PRICR APPLICATION NUMBER: US 09/301,085
PRICR FILING DATE: 199-04-28
PRICR FILING DATE: 199-04-28
PRICR FILING DATE: 199-04-28
PRICR FILING DATE: 199-04-28
PRICR FILING DATE: 1994-09-22
PRICR FILING DATE: 1994-04-13
NUMBER OF SEQ ID NOS: 214
SOFTWARE: FASTESQ for Windows Version 4.0
SEQ ID NO 199
Query Match
Best Local Similarity
"heres 2; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TYPE: PRT ; ORGANISM: Arabidopsis thaliana US-10-613-765-199
                                                                                                     ; ORGANISM: Homo sapiens US-09-792-686A-34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-792-686A-34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-10-613-765-199
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                                                                                                                                                                                FILE REFERENCE: 07039-250001
CURRENT APPLICATION NUMBER: US/09/792,686A
CURRENT FILING DATE: 2001-02-23
NUMBER OF SEQ ID NOS: 37
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 34
                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 34, Application US/09792686A Publication No. US20030068645A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 199, Application US/10613765
Publication No. US20040172673A1
GENERAL INFORMATION:
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                          APPLICANT: Goronzy, Jorg J.
APPLICANT: Kopecky, Stephen L.
APPLICANT: Weyand, Cornelia M.
TITLE OF INVENTION: METHODS AND MATERIALS FOR EVALUATING
TITLE OF INVENTION: CARDIOVASCULAR CONDITIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: RPS2 GENE FAMILY, PRIMERS, PROBES, AND TITLE OF INVENTION: DETECTION METHODS FILE REFERENCE: 00786/254005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: AUSUBEL, FREDERICK M. APPLICANT: STASKAWICZ, BRIAN J. APPLICANT: KATAGIRI, FUMIAKI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
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                                                                                                                                              LENGTH: 5
TYPE: PRT
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nes 2; Conserv
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ELLIS, JEFFREY
SALMERON, JOHN
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KUNKEL, BARBARA N.
MINDRINOS, MICHAEI
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                  86.7%; Score 13; DB 10; ilarity 66.7%; Pred. No. 1.4e+06; Conservative 0; Mismatches 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 13; DB 16; Length 4; Pred. No. 1.4e+06;
                                                          DB 10; Length 5;
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                      Gaps
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YXO YXO

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APPLICANT: TRUNEH, ALEMSEGED
APPLICANT: WANTAM, TREVOR A.
TITLE OF INVENTION: ANTI-RANK LIGAND MONOCLONAL ANTIBODIES
TITLE OF INVENTION: USEFUL IN TREATMENT OF RANK LIGAND MEDIAT
FILE REFERENCE: GP50033
CURRENT APPLICATION NUMBER: US/10/344,779
CURRENT FILLING DATE: 2003-02-17
PRIOR APPLICATION NUMBER: US/10/26161
PRIOR APPLICATION NUMBER: US 60/26,524
PRIOR APPLICATION NUMBER: US 60/230,639
PRIOR FILING DATE: 2000-09-07
NUMBER: OS 580 ID NOS: 18
PRIOR FILING DATE: 2000-09-07
NUMBER: OS 580 ID NOS: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-376-121A-143; Sequence 143, Ap.; Publication No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; LENGTH: 5
; TYPE: PRT
; ORGANISM: Mus musculus
US-10-344-779-8
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: SWEET APPLICANT: TORN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: Harley,
                                                                                        COUNTRY: USA
ZIF: 30309-3400
COMPUTER READABLE FORM:
MEDIUM TYPE: Flopby disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA: US/10/376,121A
FILING DATE: 27-Mar-2003
CLASSIFICATION: cUnknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: METHODS AND REAGENTS FOR DIAGNOSIS OF AUTOANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 218
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      APPLICATION NUMBER: 07/867,819
FILING DATE: April 13, 1992
APPLICATION NUMBER: 07/648,205
FILING DATE: January 31, 1991
                                                                                                                                                                                                                                                                                                                                                              CITY: Atlanta
STATE: GA
                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Patrea L. Pabst
STREET: Suite 2000, 1201 West Peachtree Street,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Application US/10376121A
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Pred. No. 1.4e+06;
0; Mismatches 1; Indels
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                                                                                        RESULT 9
US-10-394-471B-3
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Sequence 3, Application US/10394471B
Publication No. US20040185047A1
GENERAL INFORMATION:
APPLICANT: Giles-Komar, Jill; Carton, Jill; Scallon, Bernard
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; ORGANISM: Mus musculus US-10-738-809-8
                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/738,809
CURRENT FILING DATE: 2003-12-17
PRIOR APPLICATION NUMBER: US/10/344,779
PRIOR FILING DATE: 2003-02-19
PRIOR APPLICATION NUMBER: PCT/US01/26161
PRIOR FILING DATE: 2001-08-21
PRIOR APPLICATION NUMBER: US 60/226,524
PRIOR APPLICATION NUMBER: US 60/230,639
PRIOR APPLICATION NUMBER: US 60/230,639
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-10-376-121A-143
           Query Match
Best Local Similarity 65...
Conservative
                                                                                                                                                                                             PRIOR FILING DATE: 2000-09-07
NUMBER OF SEQ ID NOS: 18
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 8
LENGTH: 5
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Best Local Similarity
Matches 2; Conserv
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Publication No. US20040171117A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: SWEET, RAYMOND W.
APPLICANT: TORNETTA, MARK A.
APPLICANT: TRUNEH, ALEMSEGED
APPLICANT: WATTAM, TREVOR A.
TITLE OF INVENTION: ANTI-RANK LIGAND MONOCLONAL ANTIBODIES
TITLE OF INVENTION: USEFUL IN TREATMENT OF RANK LIGAND MEDIATED DISORDERS
TITLE REFERENCE: GP50033
                                                                                                                                                                               TYPE: PRT
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TELEPAX: (404)-817-8588
INFORMATION FOR SEQ ID NO: 143:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 07/472,947
FILING DATE: January 31, 1990
ATTORNEY/AGENT INFORMATION:
NAME: Pabst, Patrea L.
REGISTRATION NUMBER: 31,284
REFERENCE/DOCKET NUMBER: OMRF114
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: peptide SEQUENCE DESCRIPTION: SEQ ID NO: 143:
1 GXY 3
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Pred. No. 1.4e+06;
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                                                                                  Length 5;
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; SOFTWARE: PatentIn Ver 3.1; SEQ ID NO 3; LENGTH: 5; TYPE: PRT; ORGANISM: Homo sapiens US-10-394-471B-3
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US-09-990-762-36
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publication No. US20040185047A1
GENERAL INFORMATION:
APPLICANT: Giles Komar, Jill; Carton, Jill; Scallon, Bernard J.
APPLICANT: Giles Komar, Jill; Carton, Jill; Scallon, Bernard J.
APPLICANT: CENO202
FILE REFERENCE: CENO202
CURRENT APPLICATION NUMBER: US/10/394,471B
CURRENT FILING DATE: 2003-03-21
PRIOR APPLICATION NUMBER: 60/367,903
PRIOR APPLICATION NUMBER: 60/367,903
PRIOR APPLICATION NUMBER: 60/367,903
PRIOR APPLICATION NUMBER: 60/367,903
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SOFTWARE: PatentIn Ver 3.1
SEQ ID NO 11
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Best Local 9
                SEQ ID NO 36
LENGTH: 6
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GENERAL INFORMATION:
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -10-394-471B-11
                                                APPLICANT: JOUNG, J. KEITH
APPLICANT: MILLER, JEFFREY
APPLICANT: PABO, CARL O.
TITLE OF INVENTION METHODS AND COMPOSITIONS FOR INTERACTION TRAP ASSAYS
FILE REFERENCE: MTV-030.02 (20021-3002)
CURRENT APPLICATION NUMBER: US/09/990,762
CURRENT FILING DATE: 2001-11-14
PRIOR APPLICATION NUMBER: 09/858,852
PRIOR FILING DATE: 2001-05-16
PRIOR PHILING DATE: 2001-05-16
PRIOR PHILING DATE: 2000-05-16
NUMBER OF SEQ ID NOS: 97
SOFTMARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: ANTI-THE ANTIBODIES, COMPOSITIONS, METHODS AND USES FILE REFERENCE: CENO202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Homo sapiens
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ENGTH:
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2; Conserv
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Pred. No.
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Pred. No. 1.4e+06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 2; Conserve
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Patent No. US20020119498A1

GENERAL INFORMATION:
APPLICANT: JOUNG, J. KEITH
APPLICANT: MILLER, JEFEREY
APPLICANT: PABO, CARL O.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INTERACTION TRAP ASSAYS
FILE REFERENCE: MTV-030.02 (20021-3302)
CURRENT APPLICATION NUMBER: US/09/990,762
CURRENT APPLICATION NUMBER: US/09/990,762
CURRENT FILING DATE: 2001-11-14
PRIOR APPLICATION NUMBER: 09/858,852
PRIOR APPLICATION NUMBER: 09/858,852
PRIOR APPLICATION NUMBER: 60/204,509
PRIOR FILING DATE: 2000-05-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 40
                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                         SOFTWARE: PatentIn Ver. SEQ ID NO 41
                                                                                                                                                                                                                                                                                                                                                                                 Sequence 41, Application US/09990762 Patent No. US20020119498A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                               APPLICANT: JOING, J. KEITH
APPLICANT: MILLER, JEFFREY
APPLICANT: MILLER, JEFFREY
APPLICANT: PABO, CCARL O,
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INTERACTION TRAP ASSAYS
FILE REFERENCE: MTV-030.02 (20021-3002)
CURRENT FILING DATE: 2001-11-14
PRIOR APPLICATION NUMBER: 05/09/990,762
CURRENT FILING DATE: 2001-11-14
PRIOR APPLICATION NUMBER: 09/858,852
PRIOR FILING DATE: 2001-05-16
PRIOR APPLICATION NUMBER: 60/204,509
PRIOR FILING DATE: 2000-05-16
NUMBER OF SEO ID NOS: 97
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SOFTWARE: Patentin Ver. :
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TYPE: PRT
                  LENGTH: 6
TYPE: PRT
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Pred. No. 1.4e+06;
0; Mismatches 1
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Pred. No. 1.4e+06;
0; Mismatches 1
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Sequence 6, Application US/09518737

Publication No. US20030008321A1

GENERAL INFORMATION:
APPLICANT: FUKUI, YASUHISA
APPLICANT: SHIRAI, RYUICHI
FIILE OF INVENTION: MONOCLONAL ANTIBODY RECOGNIZING
TITLE OF INVENTION: MONOCLONAL ANTIBODY RECOGNIZING
FILE REFERENCE: 1965/49618

CURRENT APPLICATION NUMBER: US/09/518,737

CURRENT APPLICATION NUMBER: US/09/518,737

CURRENT FILING DATE: 1999-03-03

PRIOR APPLICATION NUMBER: JP 1999-250209

PRIOR FILING DATE: 1999-09-03

NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver. 2.1

LENGTH: 6
TYPE: PRT
ORGANISM: Mus musculus
US-09-518-737-6
                                                                                                                                                 Sequence 36, Application US/09858852A

Publication No. US20030044787A1

GENERAL INFORMATION:
APPLICANT: JUNE FEREY
APPLICANT: MILLER, JEFFREY
APPLICANT: MILLER, JEFFREY
APPLICANT: MILLER, JEFFREY
APPLICANT: MILLER, JEFFREY
APPLICANT: MITHODS AND COMPOSITIONS FOR INTERACTION TRAP ASSAYS
FILE REPERENCE: MTV-030.01 (20021-3001)
CURRENT FILING DATE: 2001-05-16
PRIOR APPLICATION NUMBER: 60/204,509
PRIOR APPLICATION NUMBER: 60/204,509
PRIOR APPLICATION NUMBER: 60/204,509
STWARE: Patentin Ver. 2.1
SOFTWARE: Patentin Ver. 2.1
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; OTHER INFORMATION: Description of Artificial Sequence: Zinc finger; OTHER INFORMATION: recognition sequence
US-09-990-762-41
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US-09-518-737-6
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Best Local Similarity 66.7%; Pred. No. 1.4e+06;
Matches 2; Conservative 0; Mismatches 1.
TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Zinc finger
OTHER INFORMATION: recognition sequence
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Query Match

86.7%;

Score 13;

DB 10;

Length 6;

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Rest Local Similarity 66.7%; Pred. No. 1.4e+06;
Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps

Qy 1 GXY 3
Db 3 GAY 5

Search completed: December 13, 2004, 23:02:40

Job time: 68.6 secs
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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Maximum DB seq length: 2000000000
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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ALIGNMENTS

Ig heavy chain V region - mouse ()Species: Mus musculus (house mouse) C;Species: Mus musculus (house mouse) C;Date: 06-Jan.194 #sequence_revision 01-Dec-1995 #text_change 05-Nov-1999 C;Accession: 836850 R;Jacob, J.; Kelsoe, G. R;Jacob, J.; Kelsoe, G. submitted to the EMBL Data Library, July 1992 R;Description: In situ studies on the primary immune response to (4-hydroxy-3-nitropher A;Description: In situ studies on the primary immune response to (4-hydroxy-3-nitropher A;Description: In situ studies on the primary immune response to (4-hydroxy-3-nitropher A;Description: In situ studies on the primary immune response to (4-hydroxy-3-nitropher A;Description: In situ studies on the primary immune response to (4-hydroxy-3-nitropher A;Description: In situ studies on the primary immune response to (4-hydroxy-3-nitropher A;Description: In situ studies on the primary immune response to (4-hydroxy-3-nitropher A;Description: In situ studies on the primary immune response to (4-hydroxy-3-nitropher A;Description: In situ studies on the primary immune response to (4-hydroxy-3-nitropher A;Description: In situ studies on the primary immune response to (4-hydroxy-3-nitropher A;Description: In situ studies on the primary immune response to (4-hydroxy-3-nitropher A;Description: In situ studies on the primary immune response to (4-hydroxy-3-nitropher A;Description: In situ studies on the primary immune response to (4-hydroxy-3-nitropher A;Description: In situ studies on the primary immune response to (4-hydroxy-3-nitropher A;Description: In situ studies on the primary immune response to (4-hydroxy-3-nitropher A;Description: In situ studies on the primary immune response to (4-hydroxy-3-nitropher A;Description: In situ studies on the primary immune response to (4-hydroxy-3-nitropher A;Description: In situ studies on the primary immune response to (4-hydroxy-3-nitropher A;Description: In situ studies on the primary immune response to (4-hydroxy-3-nitropher A;Description: In situ studies on the primary immune response to (4-hydroxy-3

A;Status: preliminary
A;Molecule type: nucleic acid
A;Molecule: 1-9 <-APC>
A;Residues: 1-9 <-APC>
A;Cross-references: EMBL:X67387; NID:g50113; PIDN:CAA47799.1; PID:e51594; PID:g1333871
C;Keywords: heterotetramer; immunoglobulin

A; Reference number: S25024 A; Accession: S36850

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T-cell receptor gamma chain (2t.23) - mouse (fragment)
(;Species: Mus musculus (house mouse)
(;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 07-May-1999
(;Accession: G41946
R;Whetsell, M.; Mosley, R.L.; Whetsell, L.; Schaefer, F.V.; Miller, K.S.; Klein, J.R.
Mol. Cell. Biol. 11, 5902-5909, 1991
A;Tille: Rearrangement and junctional-site sequence analyses of T-cell receptor gamma A;Reference number: A41946; MUID:92049316; PMID:1658619
A;Accession: G41946
  8
                                                                                                                                                                                           A;Status: preliminary; not compared with conceptual translation A;Molecule type: DNA A;Residues: 1-9 WHE>
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C;Keywords: T-cell receptor
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Local Similarity 66.7%; Pred. No. 2.8e+05;
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Pred. No. 2.8e+05;
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C/Date: 24-Aug-1990 #sequenc
C/Accession: A33099
R/Nichols, J.H.; Hager, L.P.
submitted to the Protein Seq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R;Nakano, N.; Kikutani, H.; Nishimoto, H.; Kishimoto, T.
J. Exp. Med. 173, 1091-1097, 1991
A;Title: T cell receptor V gene usage of islet beta cell-reactive T cells is not restric A;Reference number: PT0209; MUID:91217621; PMID:1902501
A;Accession: PT0214
A;Mccession: PT0214
A;Mccession: PT0214
A;Mccession: ST0214
A;Mccessio
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A; Accession: A33099
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C;Keywords: T-cell receptor
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C;Species: Mus musculus (house mouse)
C;Datcs 31-Dec-1991 #sequence_revision 31-Dec-1991 #text_change 30-May-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Experimental source: B lymphocyte C; Keywords: heterotetramer; immunog
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7. Exp. Med. 173, 395-407, 1991

A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A;Reference number: PT0222; MUID:91108337; PMID:1899102

A;Accession: PT0250

A;Molecule type: DNA

A;Residues: 1-11 <YAM>
                                                                                                                                                                                                                                                                                                                                                      A;Residues:
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                                                                                                                                                                                                                                                                                                                                                                                                           A;Status: preliminary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              :Date: 24-Aug-1990 #sequence_revision 24-Aug-1990 #text_change 09-Jun-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence Database, May 1990
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Pred. No.
                                                                                                                                                                                                                                                           Score 13; DB 2;
Pred. No. 8.8e+02;
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Pred. No. 8.
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A; Residues: 1-12 <LEV>
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A; Residues: 1-12 < LEH>
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A; Accession: S47395
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                                                                                                                                                                      Query Match
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                                                                                                                                          Local Similarity
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   5 GSY 7
                                                     1 GXY 3
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2; Conserv
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R;Lehner, P.J.
submitted to the EMBL Data Library, August 1994
A;Description: Human HLA-A0201 restricted recogn
A;Reference number: $47355
A;Accession: $47391
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J. Exp. Med. 178, 317-329, 1993
A;Title: Molecular characterization of transgene-induced immunodeficiency in B-less mic A;Reference number: PH1580; MUID:93301609; PMID:8315387
A;Accession: PH1611
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ig H chain V-D-J region (wild-type clone 341) - mouse (fragment) C;Species: Mus musculus (house mouse) C;Decies: Mus musculus (house mouse) C;Date: 02-Jun-1994 #sequence_revision 02-Jun-1994 #text_change 17-Mar-1999 C;Date: 05-Mar-1994 #sequence_revision 02-Jun-1994 #text_change 17-Mar-1999 C;Date: 05-Mar-1994 #sequence_revision 02-Jun-1994 #text_change 17-Mar-1999 C;Date: 05-Mar-1995 #sequence_revision 05-Jun-1994 #sequence_revision 05-Ju
                                                                                                                                                                                                                                        A;Experimental source: bone marrow pre-B lymphocyte C;Keywords: immunoglobulin
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C;Species: Homo sapiens (man)
C;Date: 13-Van-1995 #sequence_revision 13-Jan-1995 #text_change 05-Nov-1999
C;Accession: S47395
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A;Residues: 1-12 <LEH>
A;Cross-references: EMBL:Z35712; NID:g527519; PIDN:CAA84781.1; PID:g527520; EMBL:Z3570
C;Keywords: T-cell receptor
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Score 13; DB
Pred. No. 8.8e
0; Mismatches
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Pred. No. 8.8e
0; Mismatches
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Pred. No.
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T-cell antigen receptor VJ junction beta chain - human C, Species: Homo sapiens (man)
C, Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #te)
C, Accession: S47368
R, Lehner, P.J.
submitted to the EMBL Data Library, August 1994
A, Description: Human HLA-A0201 restricted recognition of A, Reference number: S47355
                                                                  A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-13 <LEH>
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A;Reference number: PH0746; MUID:92078846; PMID:1836010
                                                                                                                                             A;Accession: S47368
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A; Residues: 1-12 < CAS>
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R;Casanova, J.L.; Romero, P.; Widmann, C.; Kourilsky, P.; Maryanski, J. Exp. Med. 174, 1371-1383, 1991
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A;Experimental source: T lympho
C;Keywords: T-cell receptor
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C;Species: Mus musculus (house mouse)
C;Bate: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PH0790
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A;Reference number: PH0746; MUID:92078846; PMID:1836010
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A; Residues: 1-12 < CAS>
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C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PH0802
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                          Cross-references: EMBL:Z35693; NID:g527477; PIDN:CAA84762.1; PID:g527478 Keywords: T-cell receptor
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;Experimental source: T lymphocyte
;Keywords: T-cell receptor
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Best Local :
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Pred. No. 8.8e+02;
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A; Reference
A; Accession:
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unidentified 6.3/40K protein [imported] - rice (fragment)
(;Species: Oryza sativa (rice)
(;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_change 09-Jul-2004
C;Accession: PQ0700
R;Konatsu, S.; Kajiwara, H.; Hirano, H.
Theor. Appl. Genet. 86, 935-942, 1993
A;Title: A rice protein library; a data-file of rice proteins separated by two-dimension
A;Reference number: PQ0696
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PH0788
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                                                                                                                                                                                                                                                                                                                                                                                                A;Cross-references: EMBL:X60893
A;Experimental source: T lympho
                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Molecule type: mRNA
A; Residues: 1-13 < CAS>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R;Casanova, J.L.; Romero, P.; Widmann, C.; Kourilsky, P.; Maryanski, J.L.
J. Exp. Med. 174, 1371-1383, 1991
A;Title: T cell receptor genes in a series of class I major histocompatibility complex-allelic exclusion and antigen-specific repertoire.
A;Reference number: PH0746; MUID:92078846; PMID:1836010
A;Accession: PH0788
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    T-cell receptor alpha chain (E22 V-alpha-8.F3.4) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PH0788
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A;Molecule type: mRNA
A;Residues: 1-13 -C.EH>
A;Cross-references: EMBL:Z35709; NID:g527513; PIDN:CAA84778.1; PID:g527514
C;Keywords: T-cell receptor
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S47383
T-cell
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A;Description: Human HLA-A0201 restricted recognition of influenza
A;Reference number: $47355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              T-cell antigen receptor VJ junction beta chain - C;Species: Homo sapiens (man) C;Date: 06-Jan-1995 #sequence_revision 06-Jan-199 C;Accession: S47383
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RESULT 15
A59018
MUC1 enhancer binding protein 70K chain MUC1EBP-70 - human (fragment)
C.Species: Homo sapiens (man)
C.Accession: A59018
R.Abe, M.; Smith, C.J.; Larson, C.J.
submitted to the Protein Sequence Database, May 1998
A;Abescription: Involvement of "Ku-like" proteins in the transcription of MUC1/DF3, a bre
A;Reference number: A59018
A;Accession: A59018
A;Accession: A59018
A;Residues: 1-14 AABE>
A;Residues: 1-14 AABE>
A;Residues: 1-14 AABE>
A;Residues: 3-Val was also found
C;Keywords: DNA binding; heterodimer
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A;Molecule type: protein
A;Residues: 1-13 < KOM>
A;Cross-references: UNIPROT:Q7M1U1
Search completed: December 13, 2004, 22:32:59 Job time : 15.2 secs
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Maximum Match 100%
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Perfect score:
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1: uniprot_sprot:*
2: uniprot_trembl:*
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  GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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   UF06 MOUSE
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Q6LD78
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AAC39714
AAB92363
Q9GJW1
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062645
Q71VQ4
Q9T2H6
Q84098
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Q9SMV3
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Q47892 fremyella d

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P29264 prunus sero

Q719d2 homo sapien

Q6jdk5 canis famil

Q98mv3 arabidopsis

Q61c19 homo sapien

Q21vq4 homo sapien

Q912h6 spinacia ol

Q84098 influenza a

Q712n1 torpedo mar

Aac39714 homo sapi

Aab92363 homo sapi

Q9jw3 lagenorhync

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	Q9r469 Q9r469 Q9r501 Q9pr501 AAD2088 Q9n6i3 Q9gn6i3 Q9r4f8 Q9r4f8 Q9r4f9 Q9r362 Q9r362 Q9r362 Q9r3017 Q9s8h8	Q9S8H8

ALIGNMENTS

RESULT 1
UF06 MOUSE
STANDARD; PRT; 8 AA.
ID UF06 MOUSE STANDARD; PRT; 8 AA.
AC P38644;
DT 01-OCT-1994 (Rel. 30, Created)
DT 01-OCT-1994 (Rel. 30, Last sequence update)
DT 05-JUL-2004 (Rel. 44, Last annotation update)
DT 05-JUL-2004 (Rel. 44, Last annotation update)
DE Unknown protein from 2D-PAGE of fibroblasts (P50) (Fragment).
OS Mus musculus (Mouse).

RESULT 2 Q9QWT0 ID Q9QWT0 PRELIMINARY; PRT; 9 AA. AC Q9QWT0; DT 01-MAY-2000 (TrEMBLrel. 13, Created) DT 01-MAY-2000 (TrEMBLrel. 13, Last sequence update) DT 01-DC-2001 (TrEMBLrel. 19, Last annotation update) DE Proteinase 3 (Myeloblastin) (Fragment). GN Name=Prtn3;	1 GXY 3	Query Match 86.7%; Score 13; DB 1; Length 8; Best Local Similarity 66.7%; Pred. No. 1.8e+06; Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;	ID UF06 MOUSE STANDARD; PRT; 8 AA. AC 93644; DT 01-OCT-1994 (Rel. 30, Created) DT 01-OCT-1994 (Rel. 30, Last sequence update) DT 05-UTL-2004 (Rel. 44, Last annotation update) DT 05-UTL-2004 (Rel. 44, Last annotation update) DE Unknown protein from 2D-PAGE of fibroblasts (P50) (Fragment). OS Mus musculus (Mouse) Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. OX NCBI_TaxID=10090; RN (1) RP SEQUENCE. RC TISSUE=Fibroblast; RC TISSUE=Fibroblast; RA MEDLINE=95009907; PubMed=7523108; RA MEDLINE=9500907; PubMed=9523108; RA MEDLINE=9500907; PubMed=

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                                                                                                                                                                                                                                     EMBL; AY203963; AAO63472.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-MAR-2004 (TrEMBirel. 27, Created)
02-MAR-2004 (TrEMBirel. 27, Last sequence update)
02-MAR-2004 (TrEMBirel. 27, Last annotation update)
Type II hair keratin (Fragment).
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Mammalia; Eutheria; I
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01-0CT-2003 (TrEMBLrel. 25, Created)
01-0CT-2003 (TrEMBLrel. 25, Last sequence up
01-0CT-2003 (TrEMBLrel. 25, Last annotation
                                                                                                                                                                                                                                                                                                             EMBL; AY328516; AAP94210.1; NON_TER 1 1
                                                                                                                                                                                                                                                                                                                                                             Liu Y., He X., Xu L., Zeng Y.; Submitted (JUN-2003) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIINE=96326114; PubMed=8703476;
Cooper J.A.Jr., Culbreth R.R.;
Looper J.A.Jr., Culbreth R.R.;
Looper J.A.Jr., Looper J. Inhibitor peptide harvested human bronchial lavage: homology to influenza A mucleoprotein.
Am. J. Respir. Cell Mol. Biol. 15:207-215(1996).
Go; Go:00010236; P:anti-inflammatory response; NAS.
SEQUENCE 11 AA; 1262 MW; 951A1C3279C9DB45 CRC64;
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Mammalia; Eutheria;
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-UN-2003 (TrEMBLrel. 24, Last annotation update)
01-UN-2003 (TrEMBLrel. 24, Last annotation update)
Neutrophil inhibitor peptide, NIP-POLYMORPHONUCLEAR neutrophil
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Homo sapiens (Human).
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                                                                                                                                                                                                                                                                                    1208 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chordata;
Primates;
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Pred. No. 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                 BB
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7.2e+03;
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10-OCT-2003
10-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-JUN-2000 (TrEMBLrel. 14, Created)
01-JUN-2000 (TrEMBLrel. 14, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
01-JUN-2003 (TrEMBLrel. 25, Last annotation update)
32 kDa cell wall protein (Fragment)
Nicotiana tabacum (Common tobacco).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta
Spermatophyta; Magnoliophyta; eudicotyledons; core
lamiids; Solanales; Solanaceae; Nicotiana.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=cv. PETIT HAVANA;
Blee K.A., Bonham V.A., Mitchell G.P.,
Wojtaszek P., Bolwell G.P.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Direct protein sequencing; Hormone; Neuropeptide SEQUENCE 13 AA; 1562 MW; 0240A4504B8A632B CR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Isolation, identification and localization of a second beetle antidiuretic peptide.";
Peptides 24:27-34(2003).
-!- FUNCTION: Inhibitor of fluid secretion by Malpighian tubulous second messenger. May function as an antidiuretic length of the compassion of the comp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eigenheer R.A., Wiehart U.M., Nicolson S.W., Hull J.J., Schooley D.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota; Neoptera; Endopterygota; Coleoptera; Polyphaga; Cucujiformia; Tenebrionidae; Tenebrio.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-OCT-2003 (Rel. 42, Created)
10-OCT-2003 (Rel. 42, Last sequence update)
05-UUL-2004 (Rel. 44, Last annotation update)
Antidiuretic factor B (ADFb).
Tenebrio molitor (Yellow mealworm).
                                                                                                                                                                                 Cell wall.
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                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                          Planta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE.
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                                                                                                                                                                                                                                                                                                                                                                                                                                         "Proteomic study of secondary cell wall
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=4097;
                                                                                                                                                                                                                                                                                           SUBCELLULAR LOCATION: CELL WALL. TISSUE SPECIFICITY: XYLEM.
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                                                                                                                                                                                                                                                        GO:0005618; C:cell wall; IEA.
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Pred. No.
                                   Score 13; DB 2;
Pred. No. 8.3e+03;
                                                                                                                                             D5022297D8697DC6 CRC64;
   Mismatches
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Q7M2U2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Wachter A., Steininger H., Rausch T., Bogs J.; Submitted (MAY-2003) to the EMBL/GenBank/DDBJ databases. EMBL, AJ564376; CAD91995.1; ... GO; GO:0004357; F:Glutamate-cysteine ligase activity; IEA. GO; GO:0016874; F:ligase activity; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosid
eurosids II; Brassicales; Brassicaceae; Brassica.
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Gamma-glutamylcysteine synthetase (EC 6.3.2.2) (Fragment).
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"A rice protein library; a data-file of rice proteins separated by two-dimensional electrophoresis.";
Theor. Appl. Genet. 86:935-942(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
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Pred. No. 8.3e+03;
0; Mismatches 1;
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Pred. No. 8.3e+03;
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RESULT 12
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DT 01-0C
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DT 02-Unkno
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OC Eukar
OC Eukar
OC Sperm
OC PACCP
OX NCB1-
RN [1]
RP SEQUE
RA TOURE
RA MAIZE
DR MAIZE
CRW Direc
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Matches 2
   Query Match
Best Local Similarity
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                                                                                                                               genome analysis program.";
Theor. Appl. Genet. 93:997-1005(1996).
-i- MISCELLANEOUS: On the 2D-gel the dete
protein is: 5.9, its MW is: 34.6 kDa.
Maize-2DPAGB; P80620; COLEOPTILE.
                                                     SEQUENCE
                                                                     NON TER
                                                                                                                                                                                                             Touzet P., Riccardi F., Morin C., Damerval C., Huet J.-C., Pernollet J.-C., Zivy M., de Vienne D., "The maize two dimensional gel protein database: towards a
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"Prunus serotina amygdalin hydrolase and prunasin hydrolase.";
Plant Physiol. 100:282-299(1992).
-i- CATALYTIC ACTIVITY: (R)-prunasin + H(2)O = mandelonitrile
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids I; Rosales; Rosaceae; Amygdaloideae; Prunus.
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P29265;
P19265;
P1920 (Rel. 24, Created)
O1-DEC-1992 (Rel. 24, Last sequence update)
O5-JUL-2004 (Rel. 44, Last amnotation update)
Prunasin beta-glucosidase IIB (EC 3.2.1.118) (Prunasin hydrolase isozyme IIB) (PH_IIB) (Fragment).
                                                                                                                                                                                                                                                                                                                               Spermatophyta; Magnoliophyta; Liliopsida; Poal PACCAD clade; Panicoideae; Andropogoneae; Zea.
                                                                                                   Direct protein sequencing.
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                                                                                                                                                                                                                                                                                                                                                                                                 (Fragment)
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SUBUNIT: Monomer.
DEVELOPMENTAL STAGE: Absent from maturing black cherry fruits
DEVELOPMENTAL STAGE: Absent from maturing black cherry fruits
until 6 weeks after flowering. Then, concemitant with cotyledc
development, the level of enzyme increases with specificity for
                                                                                                                                                                                                                                                                                                                    TaxID=4577;
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2; Conserv
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yta; Magnoliophyta; Liliopsida; Poales; Poaceae;
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Pred. No. 9.4e
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Score 13; DB 1; Pred. No. 9.4e+03;
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RESULT 15
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Q9S8L4;
01-MAY-2000
  Q47893
                                                                                                                                                                                                                      SEQUENCE FROM N.A.

MEDLINE=86233345; PubMed=3086870;

Conley P.B., Lemaux P.G., Lomax T.L., Grossman A.R.;

Conley P.B., Lemayor Light harvesting polypeptides are clustered the genome of the cyanobacterium Fremyella diplosiphon.";

Proc. Natl. Acad. Sci. U.S.A. 83:3924-3928(1986).
                                                                                                                                                                                                                                                                                                                                01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-UN-2003 (TrEMBLrel. 24, Last annotation update)
Beta-allophycocyanin (Fragment).
Fremyella diplosiphon (Calothrix PCC 7601).
Bacteria; Cyanobacteria; Nostocales; Microchaetaceae; Fremyella.
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01-MAY-2000 (TIEMBLIEL 13, Last sequence update)
01-JUN-2000 (TIEMBLIEL 13, Last sequence update)
01-JUN-2000 (TIEMBLIEL 14, Last sannotation update)
Beta-D-glucoside glucohydrolase (EC 3.2.1.21) (Fragment).
Zea mays (Maize).
Zea mays (Maize).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Spermatophyta; Magnoliophyta; Andropogoneae; Zea.
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01-NOV-1996
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GO: 000422; F:beta-glucosidase activity; IEA. (SEQUENCE 15 AA; 1667 MW; 46C40370DC8C730B CRC64;
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AC Q47833;
DT 01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
DT 01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
DT 01-NOV-1998 (TrEMBLrel. 08, Last annotation update)
DE Beta-phycocyanin (Fragment).

OS Fremyella diplosiphon (Galothrix PCC 7601).

OC Bacteria; Cyanobacteria; Noscocales; Microchaetaceae; Fremyella.

OX NCBI_TaxID=1197;
RN [1]

RP SEQUENCE FROM N.A.

RX MEDILNE=86233345; PubMed=3086870;
RA Conley P.B., Lemaux P.G., Lomax T.L., Grossman A.B.;
RO Conley P.B., Lemaux P.G., Lomax T.L., Grossman A.B.;
RT "Genes encoding major light-harvesting polypeptides are clustered on "Genes encoding major light-harvesting polypeptides are clustered on "Genes encoding major light-harvesting polypeptides are clustered on "FT "Genes encoding major light-harvesting polypeptides are clustered on "FT "Genes encoding major light-harvesting polypeptides are clustered on "FT "Genes encoding major light-harvesting polypeptides are clustered on "FT "Genes encoding major light-harvesting polypeptides are clustered on "FT "Genes encoding major light-harvesting polypeptides are clustered on "FT "Genes encoding major light-harvesting polypeptides are clustered on "FT "Genes encoding major light-harvesting polypeptides are clustered on "FT "Genes encoding major light-harvesting polypeptides are clustered on "FT "Genes encoding major light-harvesting polypeptides are clustered on "FT "Genes encoding major light-harvesting polypeptides are clustered on "FT "Genes encoding major light-harvesting polypeptides are clustered on "FT "Genes encoding major light-harvesting polypeptides are clustered on "FT "Genes encoding major light-harvesting polypeptides are clustered on "FT "Genes encoding major light-harvesting polypeptides are clustered on "FT "Genes encoding major light-harvesting polypeptides are clustered on "FT "Genes encoding major light-harvesting polypeptides are clustered on "FT "Genes encoding major light-harvesting polypeptides are clustered on "FT "Genes encoding polypeptides are clustered on "Genes encoding polypep
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 number of hits satisfying chosen parameters:
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1: geneseqp1980s:*
2: geneseqp1990s:*
3: geneseqp2000s:*
4: geneseqp2001s:*
5: geneseqp2003as:*
6: geneseqp2003bs:*
7: geneseqp2003bs:*
8: geneseqp2004s:*
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             US-10-068-905-9
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Copyright (c) 1993 - 2004 Compugen Ltd.
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AAR24318
AAR24318
AAU05432
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AAU05433
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AAU05
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Abg61618 Tetrapept
Abg67985 Human CD6
Abg67985 Human CD6
Abg67987 Human SF1
Acc84564 Mutant ph
Aap71550 ADIS viru
Aap71555 AIDS viru
Aap71555 AIDS viru
Aap71555 AIDS viru
Aap83009 Blocking
Aap83011 Blocking
Aap83011 Blocking
Aap830185 Pentapept
Aar26901 Herpes si
Aar47831 Peptide T
Aar47331 Peptide T
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Aar59492 Peptide T
Aar61487 Peptide T
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Aau05432 Synthetic
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   RESULT 1
AAR44341
ID AAR44
XX AAR4
AC 
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14-MAY-1992;
17-JUL-1992;
09-DEC-1992;
                          New linear and cyclic peptide T analogues - used for treatment and prevention of inflammation, e.g. multiple sclerosis and myelopathies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14-OCT-1993.
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22-JUN-1994
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F	л 0 л 1	3	Aar44175	Adp70014	Ad198797	7		_	w	Abp67974	٠,	p67973	Abp53799	96963	Aab70331	Aab99512	Aau05467	Aay83308	Aar98997	Aar61485
10 10		۳.	Snake ven	Skin bind	Human leu	AIDS viru		≰.	≴.	Human CD6	O	Human CD6	Activin r	Human leu	Osteoclas	(D)	ct	eptide	Peptide T	Peptide T

ALIGNMENTS

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Inflammation; antiinflammatory; immunomodulator; HTLV-1; multiple sclerosis; myopathy; chronic fatigue syndrome; toxic shock; arthritis; inflammatory bowel disease; host-versus-graft; graft-versus-host; transplant.
WPI; 1993-336832/42.
                               Andersen AJ, Aston R,
Phipps DJ, Rathjen D,
                                                                                                                                                                                                                                                                                                                                                                  Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Peptide T related peptide for treating inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAR44341 standard; peptide; 4 AA.
                                                                                       (PEPT-) PEPTIDE TECHNOLOGY LTD (DRUG-) DRUG ROYALTY CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (revised)
(first entry)
                                                                                                                                        92US-00858832.
92DK-00000645.
92US-00915118.
92US-00987674.
                                                                                                                                                                                                                                     93WO-GB000649
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1. .4
/note= "one or more
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XX AAAE5
XX AAAE5
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26-NOV-1995
  The invention relates to a new method of treating eczema or dermatitis using Peptide T and its variants. The present sequence represents one
                                                                                                                                     Treatment or prevention of eczema or dermatitis, espusing specified peptide, e.g the octa:peptide Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Misc-difference
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                                                                                     Claim 3; Page 15; 20pp; English.
                                                                                                                                                                                                                              WPI; 1995-139385/18.
                                                                                                                                                                                                                                                                                Michaelis J,
                                                                                                                                                                                                                                                                                                                                                                                              24-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-SEP-1994;
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                                                                                                                                                                                                                                                                                                                                        (PEPT-) PEPTIDE TECHNOLOGY LTD
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Pred. No.
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RESULT 3
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                                                                                                                                                                                                                                                                                                                                                              The invention relates to a new method of treating disease and/or ulcerative colltis using peptide T present sequence is one of the preferred peptides. 2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Treatment or prevention of Crohn's disease or ulcerative colitis - by use of specified peptide, e.g. Peptide {\tt T.}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1995-139386/18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Michaelis J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Misc-difference
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26-NOV-1995
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AAB24318 standard; peptide; 4 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 3; Page 15; 24pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Misc-difference
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                                                                                                                                                                      1 TXY 3
                                                                                                                                                                                                               Similarity 2; Conserv
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                                                                                                                       1 TTY 3
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2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    prototype; octapeptide; Crohn's disease; ulcerative; colitis.
                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sleigh MJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       85.7%;
                                                                                                                                                                                                                                            85.7%;
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Pred. No. 1.7e
0; Mismatches
                                                                                                                                                                                                                    Score 12; DB
Pred. No. 1.76
0; Mismatches
                                                                                                                                                                                                                                    DB 2; L
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1.7e+06;
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                                                                                                                                                                                                                                                                      Length 4;
                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                        or preventing Crohn' and its variants. Th . (Updated on 25-MAR-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
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RESULT 5
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Best Local &
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           therapy. A pharmaceutical composition or vaccine production and in gene be used to treat a patient with cancer, especially prostate cancer, the vaccine can also be used to inhibit the development or progression of cancer. The polypeptides and polymcleotides can be used to diagnose cancers, especially prostate cancer. A transgenic animal comprising 24P4C12 can be used for the development and screening of therapeutic reagents. The polypeptide is a transmembrane protein which is expressed specifically in prostate cancer, allowing the development of more specific anticancer therapies and diagnostic assays.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 designated 24P4C12, and its encoded protein. 24P4C12 has anticance cytostatic activity, and can be used in vaccine production and in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes a pudesignated 24P4C12, and its encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel 24P4C12 polypeptides and polynucleotides, used treatment of cancer, especially prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; prostate tumour transmembrane protein;
                                                                                                              Tetrapeptide
Multivariate
                                                                                                                                                                        Synthetic
                                                                                                                                                                                                           24-OCT-2001
                                                                                                                                                                                                                                                  AAU05432;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 4 AA;
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                                      WO200136980-A2
                                                                         Synthetic
                                                                                                                                                                                                                                                                                      AAU05432 standard; peptide;
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2; Conserv
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                                                                                                                                                                    tetrapeptide ligand
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 35; 137pp; English.
                                                                                                            ligand; drug design; high throughput screening; HTS; MQSAR; Quantitative Structure Activity relationships.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        describes a prostate tumour associated gene, nd its encoded protein. 24P4Cl2 has anticancer and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Leong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              associated diagnosis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 12; DB
Pred. No. 1.7e
0; Mismatches
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anticancer; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 3; L...
1.7e+06;
1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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ABG61618
ID ABG6
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Best Local
                                                                                                                                                 Human; serine protease; dipeptidyl peptidase IV-related protein; DPRP; DPPIV; infection; human immunodeficiency virus; HJV-1; HIV-2; pain; diabetes; infertility; obesity, anorexia; Parkinsnon's disease; stroke; heart failure; hypertension; urinary retention; osteoporosis; cancer; ulcer; allergy; cancer; psychotic disorder; neurological disorder; ulcer; allergy; cancer; psychotic disorder; neurological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 4 AA;
                                                                       Unidentified
                                                                                                                                                                                                                                                                  Tetrapeptide amide inhibitor #3 used to inhibit DPPIV activity.
                                                                                                                                                                                                                                                                                                                                                ABG61618;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-432565/46
                                                                                                                                   dyskinesia; reproductive disorder;
                                                                                                                                                                                                                                                                                                         12-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                   ABG61618 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lundstedt T,
                                                                                                              netabolic
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2; Conserv
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                                                                                                              disorder.
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25-MAY-2001

18-APR-2002

inflammatory

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cc 7) fractional factorial design +3 cp (or 1 cp 2 random), which were used to model the interaction of a ligand with its target. The invention crelates to characterising the interaction between a Ligand Y and a Target X by obtaining information (e.g Multivariate Quantitative Structure Activity relationships, MOSAR) representing one or more physical and/or chemical properties of type X and type Y to produce a model of interaction. The methods of the invention are useful for identifying coutliers of type X or outliers of type X, drug design, design or cidentification of lead compounds (e.g by high throughput screening, HTS), design of ligands of type Y with improved affinity and/or selectivity for targets of type X, protein engineering, design of LNA or RNA molecules, consign of artificial targets of type X/or artificial ligands of type Y, analysis and/or in the engineering of regions and/or parts of targets of type X and/or ligands of type Y, design of binding a molecule, peptidomemetic, protein, enzyme, antibody, molecule, macromolecule, DNA, or a carbohydrate. The methods are also useful for designing new
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A novel method for identifying the interaction site, binding site or active site in a macromolecule, using of informative combinatorial chemistry, informative peptide libraries and Multivariate Quantitative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 5; Fig 27; 131pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Structure Activity Relationships.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-NOV-2000; 2000WO-GB004420.
                                                                                                                                                                                                                                     ligands for known targets and/or for new targets
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MELA-) MELACURE THERAPEUTICS (PETT/) PETT C P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequence represents a tetrapeptide ligand selected according to 2 (12 fractional factorial design +3 cp (or 1 cp 2 random), which were used
    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99GB-00027346
                                         85.7%;
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Score 12; DB 4;
Pred. No. 1.7e+06;
D; Mismatches 1
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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to the isolation of novel human serine proteases referred to as dipeptidyl peptidase IV (DPPIV) -related proteins (DPRP) -rhe dipeptidyl peptidase IV-related proteins (DPRP) and nucleic acids encoding them are useful for treating infections such as fungal, bacterial, protozoan and viral infections, particularly infections caused by human immunodeficiency virus (HIV-1) pain, diabetes, precocious puberty, infertility, obesity, anorexia, bulimia, parkinson's disease, acute heart failure, hypotension, hypotension, urinary retention, osteoporosis, angina pectoris, stroke, ulcers, asthma, allergies, cancers, migraine, vomiting, psychotic and neurological disorders (e.g. anxiery, dementia, or schizophrenia), and dyskinesias. These may also be used in discovering therapeutic agents for the treatment of reproductive, inflammatory and metabolic disorders. ABG61616 ABG61620 represent tetrapeptide amide inhibitors used to inhibit DPPIV
                                                                                                                                                                                                                               Human; CD66; CEACAM; cytostatic; antiinflammatory; immunomodulator; antibacterial; virucide; gene therapy; vaccine; neutrophil; immune system; autoimmune disease; cancer; infection; bacterial; virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New dipeptidyl peptidase IV-related proteins and nucleic acids encoding the proteins, useful for treating e.g. fungal, bacterial, protozoan and viral infections, cancers, allergies, neurological disorders, or pain.
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    Skubitz
                                                                       28-FEB-2001; 2001US-0272113P
                                                                                                     27-FEB-2002; 2002WO-US005720
                                                                                                                                06-SEP-2002
                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                        Human CD66 family modulating peptide SEQ ID NO
                                                                                                                                                                                                                                                                                                                     10-DEC-2002
                                                                                                                                                                                                                                                                                                                                                  ABP67985;
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                              (SKUB/)
                                                                                                                                                            WO200268601-A2
                                                                                                                                                                                                                 inflammatory disease; transplantation;
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XX O6-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to an isolated peptide (I) from a surface exposed region of a CD66 family member (ABP67130-ABP67990). The peptide modulates: (a) activation of neutrophils; (b) activation or inhibition of T-cells, B-cells, NK cells, LAK cells, dendritic cells or other immune system cells; (c) proliferation and/or differentiation of the above cells, including epithelial cells; (d) homotypic and/or heterotypic adhesion among CD66 family members; and (e) adhesion of CD66 family members; and (e) adhesion of CD66 family members and/or their ligands, such as activation of neutrophils and activation, inhibition, proliferation and/or differentiation of the immune cells. The peptides may also be used in treating or diagnosing autoimmune diseases, cancer, infections (e.g. bacterial or viral) or inflammatory diseases, in transplantation
                                                                                                                                                                                                                                                                                                                                                                     Human; CD66; CEACAM; cytostatic; antiinflammatory; immunomodulator; antibacterial; virucide; gene therapy; vaccine; neutrophil; immune eystem; autoimmune disease; cancer; infection; bacterial; virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Human CD66 family modulating peptide SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New peptide from a surface exposed region of a CD66 family member, useful for modulating the function of CD66 family members, e.g. activation of neutrophils, for treating or diagnosing autoimmune diseases or cancer, and as a vaccine.
                                                                                                                                                     (SKUB/)
                                                                                                                                                                                                                                                                                                                            Homo sapiens.
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                                                                                                                        Skubitz KM,
                                                                                                                                                                                                  28-FEB-2001; 2001US-0272113P
                                                                                                                                                                                                                                                                06-SEP-2002.
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                                                                                                                                                                                                                                 27-FEB-2002; 2002WO-US005720
                                                                                                                                                                                                                                                                                                                                                        inflammatory disease; transplantation;
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                                                                                                                        Skubitz APN
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66.7%;
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New peptide from a surface exposed region of a CD66 family member, useful for modulating the function of CD66 family members, e.g. activation of neutrophils, for treating or diagnosing autoimmune diseases or cancer, and as a vaccine.

WPI; 2002-706981/76

Claim

22

Page

47; 96pp;

English

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RESULT 9
AAE38078
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19-AUG-2002;
06-DEC-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to an isolated peptide (I) from a surface exposed region of a CD66 family member (ABP67130-ABP67990). The peptide modulates: (a) activation of neutrophils, (b) activation or inhibition of T-cells, B-cells, IAK cells, dendritic cells or other immune system cells; (c) proliferation and/or differentiation of the above cells, including epithelial cells; (d) homotypic and/or heterotypic adhesion among CD66 family members; and (e) adhesion of CD66 family members; and (e) adhesion of CD66 family members to other ligands. The peptide is useful in modulating the function of CD66 family members and/or their ligands, such as activation of neutrophils and activation, inhibition, proliferation and/or differentiation of the immune cells. The peptides may also be used in treating or diagnosing autoimmune diseases, cancer, infections (e.g. bacterial or viral) or inflammatory diseases, in transplantation
                                                                   New human monoclonal antibody that binds to human CD30, useful for treating or preventing tumor or autoimmune disease, e.g., rheumato
                                                                                                                                                                                                                                                                                                                                                                                                                     Human; antibody; CD30; tumour; autoimmune disease; rheumatoid arthritis; systemic lupus erythematosus; systemic sclerosis; Grave's disease; ALCL; atopic dermatitis; Hashimoto's thyroidittis; Chronic renal failure; AILD; acute infectious mononucleosis; angioimmunoblastic lymphadenopathy; HIV; HOdgkin's disease; Castleman's disease; Kaposi's sarcoma; lymphoma; ATL; adult T cell lymphoma; human immunodeficiency virus; carcinoma; therapy; Wegner's granulomatosis; anaplastic large cell lymphoma; Omen's syndrome; beavy chain variable domain; VH; complementarity determining region; CDR.
                                                                                                                                                                       Keler T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-NOV-2003
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                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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2002US-0404427P.
2002US-0431684P.
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Pred. No. 1.7e+06;
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The invention

relates to

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monoclonal antibody that binds

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Query Match Best Local S Matches 2

Similarity 2; Conser

Conservative

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Score 12; DB 7; Pred. No. 1.7e+06; D; Mismatches 1

Length

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Gaps

0

85.7%;

Sequence

Claim 11; Page 120; 122pp;

English

rheumatoid

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RESULT 10
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Best Local
                          The present invention relates to a new anti-CD22 antibody has variable light (VL) and variable heavy (VH) chains of the antibody RPB4. The ant CD22 antibody is useful for the manufacture of a medicament for detection inhibiting growth of CD22+ cancer cell. The present sequence represents a mutant phage sequence of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chronic renal failure, acute infectious mononucleosis, herpes or HIV (human immunodeficiency virus) virus-associated diseases. The antibody is also useful for treating Hodgkin's disease, anaplastic large cell lymphoma (AICL), adult T cell lymphoma (AIL), angioimmunoblastic lymphoma (AILCL), adult T cell lymphoma, HIV associated body cavity lymphomas, embryonal carcinomas, undifferentiated carcinomas of the pased lymphomas, embryonal carcinomas, undifferentiated carcinomas of the rhino-pharyux (e.g. Schminoke's tumour), Castleman's disease, Kaposi's Sarcoma and other T-cell or B-cell lymphomas. The present sequence is human CD30 antibody VH (heavy chain variable domain) CDR (complementarity
                                                                                                                         Example 1;
                                                                                                                                                     New anti-CD22 antibody, useful detecting or inhibiting growth heavy and light chains of RFB4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CD30. The antibody is useful for treating or preventing tumour or autoimmune disease e.g. rheumatoid arthritis, systemic lupus erythematosus, systemic sclerosis, atopic dermatitis, Grave's disease, Hashimoto's thyroiditis, Wegner's granulomatosis, Omen's syndrome, chronic renal failure, acute infectious mononucleosis, herpes or HIV
                                                                                                                                                                                                                    WPI; 2003-402972/38
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mutant phage sequence #10
                                                                                                                                                                                                                                                                                                              26-SEP-2001; 2001US-0325360P.
                                                                                                                                                                                                                                                                                                                                                                                                         WO2003027135-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cancer; phage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    anti-CD22 antibody; RFB4; Cytostatic; Gene therapy; CD22-Antagonist;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mutant;
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                                                                                                                     22; 59pp; English
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66.7%;
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Pred. No.
                                                                                                                                                     for the manufacture of a medicament for of CD22+ cancer cell, comprising variable antibody.
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                                                                                                                                                                                                                                                                                 SERVICES
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RESULT 12
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26-JUN-1986;
12-DEC-1986;
11-MAY-1987;
03-JUN-1987;
                                                                                                                                                                                                                                            This peptide is a specifically claimed example of a highly generic formula. The peptide blocks effectively receptor sites of cells and prevents cell infectivity with AIDS virus in monkey, rat and human brair membranes and cells of the immune system. Vaccine prepns. contg. the peptide provide protection against infection by AIDS virus. The peptide can also be used in kits for the detection of the AIDS virus and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-MAR-2003
07-MAY-1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AIDS; virus; receptor site; HIV; vaccine; antibodies
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       AAP71551;
                                                                                                                                                                                                 antibodies to the AIDS virus; and as immunogens to elicit monoclonal antibodies. The peptide may be produced by conventional methods of peptide synthesis using solid phase or liquid phase methods. See also AAP71548-56, and EP-249394. (Updated on 25-MAR-2003 to correct PA field.)
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                            AAP71551 standard; protein; 5
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                                                                                                                                                                                                                                                                                                                                                    New peptide(s) related to AIDS virus - which inhibit AIDS virus binding to receptor sites and prevent cell infectivity with AIDS virus.
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86US-00878586.
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87US-00048148.
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RESULT 13
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26-JUN-1986;
12-DEC-1986;
11-MAY-1987;
                                                                                                                                                                                                                                                                                                                                This peptide is a specifically claimed example of a highly generic formula. The peptide blocks effectively receptor sites of cells and prevents cell infectivity with AIDS virus in monkey, rat and human brain membranes and cells of the immune system. Vaccine prepns. conty the peptide provide protection against infection by AIDS virus. The peptide can also be used in kits for the detection of the AIDS virus and antibodies to the AIDS virus; and as immunogens to elicit monoclonal antibodies. The peptide may be produced by conventional methods of peptide synthesis using solid phase or liquid phase methods. See also AAP71548-56, and EP-249394. (Updated on 25-MAR-2003 to correct PA field.)
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07-MAY-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New peptide(s) related to AIDS virus - which inhibit AIDS virus binding to receptor sites and prevent cell infectivity with AIDS virus.
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07-MAY-1991
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               Synthetic
                                      AIDS; virus;
                                                              AIDS virus
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                                                              receptor site blocking peptide (VIII).
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86US-00878586.
86US-00940919.
87US-00048148.
87EP-00304939.
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                                      receptor site; HIV; vaccine; antibodies
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                                                                                      entry)
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66.7%;
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Pred. No. 1.7e+06;
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RESULT 14
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Best Local S
Matches 2
  03-JUN-1986;
26-JUN-1986;
12-DEC-1986;
11-MAY-1987;
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26-JUN-1986;
12-DEC-1986;
11-MAY-1987;
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                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                 AIDS; virus;
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07-MAY-1991
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                                                                                                                 03-JUN-1987;
                                                                                                                                                            16-DEC-1987
                                                                                                                                                                                                                                                                                                                                         AIDS virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAP71552 standard;
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                                                                                                                                                                                                                                                                                         receptor site; HIV; vaccine; antibodies
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(first entry)
86US-00869919.
86US-00878586.
86US-00940919.
87US-00048148.
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86US-00878586.
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Pred. No. 1.7e+06;
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RESULT 15
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Best Local S
Matches 2
                                                                                                20-AUG-1986;
01-MAY-1987;
29-JUN-1987;
07-OCT-1987;
                                                                                                                                                                              05-MAY-1988.
                                                                                                                                                                                                     GB2196634-A
                                                                                                                                                                                                                                                                         Blocking peptide used in composition infections.
                                                                                                                                                                                                                                                                                                        25-MAR-2003
10-DEC-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This peptide is a specifically claimed example of a highly generic formula. The peptide blocks effectively receptor sites of cells and prevents cell infectivity with AIDS virus in monkey, rat and human brain membranes and cells of the immune system. Vaccine prepns. contg. the peptide provide protection against infection by AIDS virus. The peptide can also be used in kits for the detection of the AIDS virus and antibodies to the AIDS virus; and as immunogens to alicit monoclonal antibodies. The peptide may be produced by conventional methods of peptide synthesis using solid phase or liquid phase methods. See also AAP71548-56, and EP-249394. (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                        19-AUG-1987;
                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                            AAP83009;
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                                                                           (GENE-)
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                                                                                                                                                                                                                                                                                                                                                               standard; protein; 5
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                                                                                                                                                                                                                                              monoclonal antibodies; neutralising region
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87US-00045026.
87US-00067996.
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Monoclonal antibodies and peptide (s) - useful of human immuno:deficiency virus infections.

for treatment and diagnosis

WPI; 1988-103268/15

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XX

This is a blocking peptide which inhibits virus proliferation. It is CC useful, in conjunction with a monoclonal antibody to a neutralising CC useful, in conjunction with a monoclonal antibody to a neutralising CC region of HIV, for treatment or prophylaxis of HIV infections. N- and C- terminals can have up to 20 amino acids attached. See also AAP33003-08 CC and AAP83010-12. This is equivalent to NLB9701950 (accession no. is from CC this). (Updated on 25-MAR-2003 to correct PF field.) (Updated on 25-MAR-2003 to correct PF field.) (Updated on 25-MAR-2003 to correct PF field.) (Updated on 25-MAR-2003 to correct DF field.) (Updated on 25-MAR-2003 to correct DF field.)

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SQ Sequence 5 AA;

Query Match
Best Local Similarity 66.7%; Pord. No. 1.7e+06;
Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Search completed: December 13, 2004, 22:22:42

Job time: 82 secs
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Result
No.
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       protein search, using
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
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      478139 seqs, 66318000 residues
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US-09-854-799-40

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US-08-672-213-33
                                                                                                          US-08-302-829-9
US-09-421-845-9
US-09-421-845-9
US-09-421-843-5
US-08-818-8408-6
US-08-403-718-8
US-08-619-462-8
US-08-619-462-8
US-09-98-2378-8
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US-09-98-231-6258-14
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US-08-302-829-9
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Patent No.
GENERAL IN
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APPLICANT: ASTON, Roger
APPLICANT: CARLEN, Peter L
APPLICANT: CARLEN, Peter L
APPLICANT: MacFADDAEN, Douglas I
APPLICANT: MHIPPS, David J
APPLICANT: RATHJEN, Deborah
APPLICANT: WIDNER, Fred
TITLE OF INVENTION: Peptide T ar
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of

d Related Peptides in Inflammation, Includi

Including Multiple Sclerosis

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Sequence 28, Appl	`	e 25,	e 1	e 400,	A	e 1162	e 1045, Ā	e 11,	equence	e 15,	11,	ω 5	, App	ent No. 5	e 121,	121,	Sequence 21, Appl	

ALIGNMENTS

9, Application US/08302829 5. 5756449

INFORMATION:

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COUNTRY: USA
ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/302,829
FILING DATE: 29-MAR-1993
CLASSIFICATION DATA:
APPLICATION NUMBER: WO PCT/GB93/00649
FILING DATE: 29-MAR-1993
PRIOR APPLICATION NUMBER: US 07/987,674
FILING DATE: 09-DEC-1992
PRIOR APPLICATION NUMBER: US 07/987,674
FILING DATE: 17-JUL-1992
PRIOR APPLICATION NUMBER: US 07/915,118
FILING DATE: 17-JUL-1992
PRIOR APPLICATION NUMBER: DX 645/92
PRIOR APPLICATION NUMBER: DX 645/92
PRIOR APPLICATION NUMBER: DX 645/92
PRIOR APPLICATION NUMBER: US 07/858,832
PRIOR APPLICATION DATA:

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CORRESPONDENCE ADDRESS:
BANNER &
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                                                  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/082,837A
FILING DATE: 21-UUN-1998
CLASSIFICATION TATA:
APPLICATION NUMBER: US 09/082,837A
FILING DATE: 21-UUN-1998
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/302,829
FILING DATE: 29-WAR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WD PCT/GB93/00649
FILING DATE: 29-WAR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WD PCT/GB93/00649
FILING DATE: 29-WAR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/987,674
FILING DATE: 09-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/987,674
FILING DATE: 09-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/915,118
FILING DATE: 17-UUL-1992
PRIOR DATE: 17-UUL-1992
PRIOR DATE: 17-UUL-1992
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INFORMATION FOR SEQ ID NO: 9:
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CITY: Chicago
TTATE: Illinois
USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: RATHJEN, Deborah
APPLICANT: WIDMER, Fred
TITLE OF INVENTION: Peptide T and Related Peptides in the Treatment of
TITLE OF INVENTION: Inflammation, Including Multiple sclerosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
                                       PRIOR APPLICATION DATA:
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PLECOMMUNICATION INFORMATION:
TELEPHONE: 312-715-1000
TELEPAX: 312-715-1234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity nes 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: Peptide
LOCATION: 1..4
OTHER INFORMATION: /note= "aal is D-Thr, aa4 is D-Thr"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
STRANDEDNESS: not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 60606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               E: BANNER & WITCOFF, LTD.
10 S. Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MacFADDAEN, Dou
PHIPPS, David J
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CARLEN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hoth
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            peptide
MBER: DK 645/92
14-MAY-1992
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66.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Douglas K
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3.8e+05;
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Best Local Similarity
Matches 2; Conserva
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OTHER INFORMATION: US-09-082-837A-9
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                                                                                                                      ; NAME/KEY: PEPTIDE; LOCATION: (1)..(4)
; LOCATION: (1)..(4)
; OTHER INFORMATION: Thr at positions 1 and 4 are both D-Thr.
US-09-421-845-9
                                                                                                                                                                                                                                                 OF INVENTION: Peptide T and Related Peptides In the Treatment of FILE REFERENCE: Peptide T Inflammation, Including Multiple Slerosis.

FILE REPERIOR: Peptide T Inflammation
CURRENT APPLICATION NUMBER: US/09/421,845
CURRENT FILING DATE: 1999-10-20
PRIOR APPLICATION NUMBER: 09/082,837
PRIOR APPLICATION NUMBER: 09/082,837
PRIOR FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 9, Application US/09421845
Patent No. 6265374
                                                            Matches
                                                                         Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -09-421-845-9
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 4 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                               LENGTH: 4
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 27-MAR-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Drehkoff, W. Dennis
REGISTRATION NUMBER: 27193
REFERENCE/DOCKET NUMBER: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: not
TOPOLOGY: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: Peptide
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                                                            Similarity
2; Conserv
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                               TXY 3
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66.7%;
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66.7%;
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Pred. No. 3.8e+05;
                                                                             Score 12; DB 3;
Pred. No. 3.8e+05;
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                                                                 Indels
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Patent No. 5534495
GENERAL INFORMATION:
                                                                                                                                                                                                                                                       Sequence 8, Applicati
Patent No. 5756449
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: 212-986-4091
TELEPAX: 212-818-9479
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/067,523
FILING DATE: 23-MAY-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 10016-2391
COMPUTER READABLE FORM:
MEDIUM TYPE: F10ppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPUTER: DOS
CORRETTING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: not relevar MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: PERT, CANDANCE B.
APPLICANT: RUFF, MICHAEL R.
TITLE OF INVENTION: TREATMENT OF NON-HIV NEUROPATHIC
TITLE OF INVENTION: PAIN SYNDROMES
CORRESPONDENCE ADDRESS:
ADDRESSEE: BANNER & ALLEGRETTI, LTD
STREET: 10 S. Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                   IITLE OF INVENTION: Peptide 	exttt{T}^{l} and Related Peptides in the IITLE OF INVENTION: Treatment, of Inflammation, Including Multiple Sclerosis NUMBER OF SEQUENCES: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: GOLDBERG, JULES E.
REGISTRATION NUMBER: 24,408
REFERENCE/DOCKET NUMBER: AD
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                                                                                                                                                                                                                                                                                               Application US/08302829
                                                                                                 ANDERSEN, Anders J
ASTON, Roger
CARLEN, Peter L
DOOB, Penelope R
MacFADDAEN, Douglas K
PHIPPS, David J
PHIPPS, David J
RATHJEN, Deborah
WIDMER, Fred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NEW YORK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SS: not relevant not relevant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          212-986-4090
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                85.7%;
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Pred. No. 3.8e+05;
0; Mismatches 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1; Indels
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Best Local S
Matches 2
                                                                                                                                                                                                                                                                Sequence 6, Application US/08481840A Patent No. 5763406
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/987,674
FILING DATE: 09-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/915,118
FILING DATE: 17-JUL-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DK 645/92
APPLICATION NUMBER: DK 645/92
                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: PEDERSEN, Ove
APPLICANT: MACCADEN, D.
TITLE OF INVENTION: Peptides for Treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 312-715-1234
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy Disk, 3.50 inch
COMPUTER: IBM PC compatible
OPERATING SYSTEM: MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                            NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Allegretti, Ltd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 94
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-715-1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/0 FILING DATE: 29-MAR-1993 CLASSIFICATION: 514
                                                                 ZIP:
                                                                                               CITY: Chicago
STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: no
TOPOLOGY: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Chicago
STATE: Illino
                                                                                                                                  STREET:
                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                     2 TSY 4
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                                                                  60606
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                                                                                                                                10 South Wacker Drive
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66.7%;
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Pred. No. 3.8e+05;
0; Mismatches 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 8, Application US/08403718 Patent No. 5795858
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                 APPLICATION NUMBER: US/08/403,711
FILING DATE: 31-MAR-1995
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: MITCHARD, LEONARD C.
REGISTRATION NUMBER: 29,009
REFERENCE/DOCKET NUMBER: 47-74
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 816-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: MICHAELIS, JURGEN
APPLICANT: SLEIGH, MERILYN J.
TITLE OF INVENTION: TREATMENT OR PREVENTION OF CROHN'S
TITLE OF INVENTION: DISEASE AND/OR ULCERATIVE COLITIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 617-345-9111 INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/4
FILING DATE: 07-JUN-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: 08/107,
FILING DATE: 24-NOV-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                    TELEFAX: (704) 816-41
                                                                                                                                                                                                                                                                                      ZIP: 22201-4/14

COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
**DOI.TCATION NUMBER: US/08/403,718
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: Linear MOLECULE TYPE: peptide
TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: NIXON & VANDERHYE P.C.
STREET: 1100 NORTH GLEBE ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
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REFERENCE/DOCKET NUMBER: 93
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: ARLINGTON
STATE: VIRGINIA
COUNTRY: U.S.A.
                                        STRANDEDNESS:
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                                                                              ENGTH:
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66.7%;
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Pred. No. 3.8e+05;
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US-08-619-462-8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 8, Application US/08619462
Patent No. 5798335
GENERAL INFORMATION:
APPLICANT: MICHAELIS, JURGEN
APPLICANT: Trigg, Timothy Elliot
APPLICANT: Trigg, Timothy Elliot
TITLE OF INVENTION: METHOD FOR THE TREATMENT OR PREVENTION OF
TITLE OF INVENTION: ECZEMA/DERMATITIS
                                                                                            Sequence 8, Application US/09082837A Patent No. 6011014
                                                                                                                                                                                                                                       Matches
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                                                     GENERAL INFORMATION:
APPLICANT: ANDERS
APPLICANT: ASTON,
                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (704) 816-4100
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 47
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 816-4000
                                         APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 1100 NON
CITY: ARLINGTON
STATE: VIRGINIA
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nes 2; Conservative
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2; Conserv
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ANDERSEN, Anders J
ASTON, Roger
CARLEN, Peter L
DOOB, Penelope R
MacFADDAEN, Douglas K
PHIPPS, David J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3: NIXON & VANDERHYE P.C
1100 NORTH GLEBE ROAD
                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                     peptide
                                                                                                                                                                                                                                      85.7%; Score 12; DB 1; Length 5; 66.7%; Pred. No. 3.8e+05; ative 0; Mismatches 1; Indels
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; Sequent No. 605/***; Fatent No. 605/***; GENERAL INFORMATION: Finothy J.; APPLICANT: Klepfer, Sharon APPLICANT: Reed, Albert Paul
                                                                                                                                               US-08-331-625A-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release: #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/082,837A
FILING DATE: 21-UN-1998
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/302,829
FILING DATE: 29-MAR-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:

APPLICATION NUMBER: DK 645/92
PILING DATE: 14-MAY-1992
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/858,8
FILING DATE: 27-MAR-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: 312-715-1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US 07/987,674
FILING DATE: 09-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/915,118
FILING DATE: 17-UU-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: LOCATOR CHICAGO CTATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: WO PCT/GB93/00649 FILING DATE: 29-MAR-1993 PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                 Local Similarity les 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: no
TOPOLOGY: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Drehkoff, W. Dennis
REGISTRATION NUMBER: 27193
REFERENCE/DOCKET NUMBER: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
                                                                                                           4, Application US/08331625A 6057436
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10 S. Wacker Drive
                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                312-715-1234
                                                                                                                                                                                                                                                                                                                                                                                                                                  not relevant
                                                                                                                                                                                                                                                                                                                 85.7%;
66.7%;
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Pred. No. 3.8e+05;
0; Mismatches 1
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                                                                                                                                                                                                                            APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                             Sequence 8, Application US/09421845 Patent No. 6265374
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Best Local (
                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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      APPLICANT: Widmer
TITLE OF INVENTION: Deptide T and Related Peptides In the TreaTITLE OF INVENTION: Inflammation, Including Multiple Slerosis FILE REFERENCE: Peptide T Inflammation
CURRENT APPLICATION NUMBER: US/09/421,845
CURRENT FILING DATE: 1999-10-20
PRIOR APPLICATION NUMBER: 09/082,837
PRIOR FILING DATE: 1998-05-21
                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (215) 270-50
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 14-NOV-1990
ATTORNEY AGENT INFORMATION:
NAME: Schreck, Patricia A.
REGISTRATION UNMBER: 33,777
REFERENCE/DOCKET NUMBER: SEC
TELECOMMUNICATION INFORMATION:
                                                                                                                                                         PPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07,
FILING DATE: 13-MAY-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/880,194
FILING DATE: 08-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/331,625A
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
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STATE: PA
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GY: unknown
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Carlen
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Rathjen
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                                                                                                                                                                                            MacFadden
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (215)
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66.7%;
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Matches Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; ORGANISM: Homo sapiens US-09-421-845-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-494-151-14
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 8
LENGTH: 5
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Patent No. 6372224

GENERAL INFORMATION:

APPLICANT: Mill
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Best Local S
Matches
                                                                   Query Match
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                                                                                                                                                                                                                                                                                                            TELEPHONE: (215) 270-5015
TELEPAX: (215) 270-5990
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/331,625
APPLICATION NUMBER: US 07/698,927
APPLICATION NUMBER: US 07/698,927
APPLICATION NUMBER: US 07/613,066
APPLICATION: DATE: 14-NOV-1990
APPLICATION: DATE
Local Similarity hes 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 19406-2799

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IEM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25
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Reed, Albert Paul
Jones, Elaine V.
TITLE OF INVENTION: Canine Coronavirus S Gene and Uses
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS
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                                                                                                                                                                            TOPOLOGY: unknown
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Schreck, Patricia A.
REGISTRATION NUMBER: 33,777
REFERENCE/DOCKET NUMBER: SBC H85010-1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 TSY 4
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2; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: SmithKline Beecham Corporation - Corporate
Patents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/09/494,151 FILING DATE: 28-Jan-2000
                                                                                                                                                                                                                                                                                  TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Application US/09494151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LING DATE: 28-Jan-2000
LASSIFICATION: <Unknown>
   Conservative
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66.7%;
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Pred. No. 3.8e+05;
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   Score 12; DB 3; Le
Pred. No. 3.8e+05;
0; Mismatches 1;
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           Indels
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TOPOLOGY: unknown

MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 14:
US-09-972-484-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 13
US-09-972-484-14
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US-09-628-665-15
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GENERAL INFORMATION:
APPLICANT: Milli
Sequence 15, Application US/09628665 Patent No. 6673771 GENERAL INFORMATION:
                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (215) 270-5
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/494,151

FILING DATE: 28-Jan-2000

APPLICATION NUMBER: 08/331,625

FILING DATE: -Unknown-
APPLICATION NUMBER: US 07/698,927

FILING DATE: 13-MAY-1991

APPLICATION NUMBER: US 07/613,066

FILING DATE: 14-NOV-1990

APPLICATION NUMBER: US 07/613,066
                                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Schreck, Patricia A.
REGISTRATION NUMBER: 33,777
REFERENCE/DOCKET NUMBER: SBC H85010-1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 270-5015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
operating system: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/972,484
FILING DATE: 05-Oct-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 19406-2799
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 59
CORRESPONDENCE ADDRESS:
ADDRESSE: SmithKline Beecham
Patents
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Jones, Elaine V.
TITLE OF INVENTION: Canine Coronavirus S Gene and Uses
Therefor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                     1 TXY 3
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CITY: King of Prussia
STATE: PA
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                                                                                                                                                                                                                                                                                                                                                        LENGTH: 5 amino acids
TYPE: amino acid
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                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                            85.7%;
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                                                                                                                                                                                                                              Score 12; DB 4; Length 5; Pred. No. 3.8e+05;
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                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                0;
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APPLICANT: Greene, Mark I.
APPLICANT: Murali, Ramachandran
APPLICANT: Kinosaki, Masahiko
ITTLE OF INVENTION: Methods of Inhibiting Osteoclast Activity
FILE REFERENCE: UPN3831
CURRENT FILING DATE: 2000-07-28
PRIOR APPLICATION NUMBER: 60/146,094
PRIOR APPLICATION NUMBER: 60/146,094
PRIOR FILING DATE: 1999-07-28
NUMBER OF SEQ ID NOS: 34
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 15
LENGTH: 5
TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
OTHER INFORMATION: Description of Artificial Sequence: No. 6673771el Sequence
                                                                                                                                                                                                                                                                                                                                      RESULT 15
5276016-4
;Patent No. 5276016
; Patent No. 5276016
; PILLEAM L.
; PARRAR, WILLIAM L.
; PARRAR, WILLIAM R.
; PAPLICATION NOWABER: US/07/568,616
; PAPLICATION NOWABER: 314,507
; PAPLICATION NOWABER: 314,507
; PAPLICATION NOWABER: 314,507
; PAPLICATION NOWABER: 48,148
; PAPLICATION NOWABER: 48,148
; PAPLICATION NOWABER: 48,148
; PAPLICATION NOWABER: 878,586
; PAPLICATION NOWABER: 869,919
; PARCATH. S.
Search completed: December 13, 2004, 22:34:52 Job time: 22.4 secs
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5276016-4
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Best Local Similarity
Matches 2; Conserv
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                                                                                                                                                                                                                     85.7%;
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Pred. No. 3.8e+05;
0; Mismatches 1; Indels
                                                                                                                                                                                           0; Gaps
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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Run
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Total number of hits satisfying chosen parameters:
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             1121005
                                                                                                                                                                                                                                                                                                                                                               Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                            Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  seq length: 0
seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                             BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            December 13, 2004, 22:31:47; Search time 68.6 Seconds (without alignments)
20.827 Million cell updates/sec
             85...7
85...7
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14
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1: /cgn2_6/ptodata/2/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/2/pubpaa/POT_NEW_PUB.pep:*

3: /cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.pep:*

4: /cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.pep:*

5: /cgn2_6/ptodata/2/pubpaa/US07_NEW_PUB.pep:*

6: /cgn2_6/ptodata/2/pubpaa/US07_NEW_PUB.pep:*
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Copyright (c) 1993 - 2004 Compugen Ltd
                                                                                                                                                                                                                                                                                                                                                                                                                                                     /cgn2_6/ptodata/2/pubpaa/US08_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US09A_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US09A_PUBCOMB.pep:*
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/cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*
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/cgn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
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                                                                                                                                                                                                                                                                          BB
14 US-10-285-045-53
14 US-10-284-660-53
15 US-10-338-366-42
9 US-09-972-484-14
US-10-181-937-57
10S-09-854-799-40
US-10-272-411-26
4 US-10-272-328A-26
4 US-10-285-045-61
14 US-10-384-660-61
15 US-10-384-669-67
15 US-10-384-569-7
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     Sequence 53, Appl
Sequence 54, Appl
Sequence 42, Appl
Sequence 14, Appl
Sequence 7, Appl
Sequence 80, Appl
Sequence 82, Appl
Sequence 26, Appl
Sequence 61, Appl
Sequence 61, Appl
Sequence 42, Appl
Sequence 42, Appl
Sequence 47, Appli
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US-10-285-045-53
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-10-461-863-8	-10-237-405-1	-10-053-498B-	-10-423-007-2	-10-053-520-22	10-022-066-23	-10-286-457-	-10-160-506-10	10-304-059-	-10-190-082-	-10-052-578-22	-10-162-004-	10-162-004-	10-162-004-7	09-954-385-4	09-996-265-14	09-996-265-1	09-996-265-14	09-996-265-14	09-996-265-8	09-996-265-8	09-992-896-2	9-185-908-12	9-996-288-14	9-996-288-14	9-996-288-14	9-996-288-14	9-996-288-8	9-996-288-8	US-09-727-889-20	R-101-2/8-1
uence 81,	equence 17,	equence 224	equence 21,	equence 224	equence 234	equence 115	equence 103	equence 9,	equence 653	equence 224	equence 9,	equence 8,	equence 7,	equence 41,	equence 149	equence 147	equence 145	equence 143	equence 85,	equence 81,	equence 29,	equence 121,	quence 149,	equence 147,	equence 145,	equence 143,	equence 85,	equence 81,	nce 20,	equence 14,
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ALIGNMENTS

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APPLICANT: Rene S. Hubert
APPLICANT: Kahan Leong
APPLICANT: Kahan Leong
APPLICANT: Douglas C. Saffran
TITLE OF INVENTION: NOVEL 13-TRANSMEMBRANE PROTEIN EXPRESSED
TITLE OF INVENTION: NOVEL 13-TRANSMEMBRANE PROTEIN EXPRESSED
TITLE OF INVENTION: NOVEL 13-TRANSMEMBRANE PROTEIN EXPRESSED
TITLE AFFERENCE: 51158-20011.10
CURRENT APPLICATION NUMBER: US/10/285,045
CURRENT APPLICATION NUMBER: US 09/547,789
PRIOR APPLICATION NUMBER: US 09/547,789
PRIOR APPLICATION NUMBER: 60/128,858
PRIOR APPLICATION NUMBER: 60/128,858
PRIOR FILING DATE: 1999-04-12
NUMBER OF SEQ ID NOS: 71
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 53
                                                                                                                                                                                  ; LENGTH: 4
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-285-045-53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 53, Application US/10285045
Publication No. US20030147904A1
GENERAL INFORMATION:
APPLICANT: Daniel E.H. Afar
APPLICANT: Rene S. Hubert
                                                                                       Query Match
Best Local Similarity
Matches 2; Conserv
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TTY 3
                                               TXY 3
                                                                                            85.7%;
ilarity 66.7%;
Conservative
                                                                                            Score 12; DB 14; ]
Pred. No. 1.4e+06;
0; Mismatches 1;
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                                                                                            Gaps
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Application US/09972484

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GENERAL INFORMATION:

APPLICANT: Daniel E.H. Afar
APPLICANT: Rene S. Hubert

APPLICANT: Kahan Leong
APPLICANT: Kahan Leong
APPLICANT: Kahan Leong
APPLICANT: Daniel E. Raitano
APPLICANT: Daniel B. Raitano
APPLICANT: Douglas C. Saffran
APPLICANT: NOUSL 13-TRANSMEMBRANE PROTEIN EXPRESSED
TITLE OF INVENTION: NOVEL 13-TRANSMEMBRANE PROTEIN EXPRESSED
TITLE OF INVENTION: NOVEL 13-TRANSMEMBRANE PROTEIN EXPRESSED
CURRENT APPLICATION NUMBER: US/10/284,660
CURRENT APPLICATION NUMBER: US 09/547,789
PRIOR APPLICATION NUMBER: 09/547,789
PRIOR FILING DATE: 2000-04-12
PRIOR APPLICATION NUMBER: 60/128,858
PRIOR APPLICATION NUMBER: 60/128,858
PRIOR APPLICATION NUMBER: 60/128,858
PRIOR FILING DATE: 1999-04-12
NUMBER OF SEQ ID NOS: 71
SOFTWARE: FASTSEQ for Windows Version 4.0
SEC ID NO 53
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Best Local Similarity
Matches 2; Conserve
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RESULT 4
US-09-972-484-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TYPE: PRT; ORGANISM: Homo sapiens US-10-284-660-53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-10-338-366-42
                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 53
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 42, Application US/10338366
Publication No. US20040006215A1
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                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Homo sapiens
-10-338-366-42
                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Trem1, John
TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES AGAINST CD30
FILE REFERENCE: MXI-180
CURRENT APPLICATION NUMBER: US/10/338,366
CURRENT FILING DATE: 2003-01-07
PRIOR APPLICATION NUMBER: US 60/347649
PRIOR FILING DATE: 2002-01-09
PRIOR FILING DATE: 2002-01-09
PRIOR FILING DATE: 2002-08-19
PRIOR FILING DATE: 2002-08-19
                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 60/431684 PRIOR FILING DATE: 2002-12-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Keler, Tib
APPLICANT: Graziano,
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Pred. No. 1.4e+06;
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Pred. No. 1
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US-10-181-937-57
                                                                                                                                        Sequence 57, Application US/10181937
Publication No. US20040055022A1
GENERAL INFORMATION:
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Patent No. U
                                                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local
               APPLICANT: Cheng, Nick
APPLICANT: Gagnier, Liane
APPLICANT: Jeffries, Wilfred A.
APPLICANT: University of British Columbia
TITLE OF INVENTION: Compositions and Methods for Screening
TITLE OF INVENTION: Agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           general information:

APPLICANT: Miller,
FILE REFERENCE: 020739-000410US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/494,151
FILING DATE: 28-Jan-2000
APPLICATION NUMBER: 08/331,625
FILING DATE: <Unknown>
APPLICATION NUMBER: US 07/698,927
FILING DATE: 13-MAY-1991
APPLICATION NUMBER: US 07/613,066
FILING DATE: 14-NOV-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/972,484
FILING DATE: 05-Oct-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Schreck, Patricia A.
REGISTRATION NUMBER: 33,777
REFERENCE/DOCKET NUMBER: SBC H85010-1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 270-5015
TELEPAX: (215) 270-5090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 19406-2799
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Canine Coronavirus S Gene and Uses
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                                                                                                                                                                                                                                                                                                                                                                          . Similarity 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 5 amino acids
TYPE: amino acid
TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 709 Swedeland Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: SmithKline Beecham Corporation - Corporate
                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Miller, Timothy J.
Klepfer, Sharon
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Reed, Albert Paul
Jones, Elaine V.
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Pred. No. 1.4e+06;
0; Mismatches 1
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Query Match
                                                                                                                                                TELEPAX: (215) 270-50:
INFORMATION FOR SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/854,799
FILING DATE: 14-May-2001
                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 08/392,459
FILING DATE: <UNKNOWN>
APPLICATION NUMBER: US 07/613,066
FILING DATE: 14-NOV-1990
ATTORNEY/AGENT INFORMATION:
                                                          TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO: 40:
                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: SBC 14532B TELECOMMUNICATION INFORMATION: TELEPHONE: (215) 270-5015
                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: SmithKling TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                     NAME: King, William T. REGISTRATION NUMBER: 30,954
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE: PA
                                                                                                                                  TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ET: 709 Swedeland
: King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SmithKline Beecham, Corporation NVENTION: Recombinant Feline Coronavirus
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Score 12;
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DB 9;
Length 6;
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/ TYPE: PRT
/ ORGANISM: Homo sapiens
US-10-272-411-26
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US-09-727-963A-82
                                                                                                                                                                                                                                                                                                                                                                                                                                          US-10-272-411-26
                                                                                                                                                SOFTWARE: PatentIn version 3.1 SEQ ID NO 26
                                                                                                                                                                                                                                                                                                                                               Sequence 26, Application US/10272411
Publication No. US20030100068A1
GENERAL INFORMATION:
APPLICANT: Barnes Jewish Hospital
APPLICANT: Lam, Jonathan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: P
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Patent No. US20020155106A1
GENERAL INFORMATION:
APPLICANT: V.I. Technologies, Inc.
APPLICANT: Hammond, David J.
TITLE OF INVENTION: METHOD OF IDENTIFYING A LIGAND FOR A TARGET MOLECULE
FILE REFERENCE: 18242-505
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 66.7
Matches 2; Conservative
 Matches
                                     Query Match
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Best Local Similarity
Matches 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                    APPLICANT: Ross, F. Patrick
APPLICANT: Teitelbaum, Steven
TITLE OF INVENTION: RANKL MIMICS AND USES THEREOF
FILE REFERENCE: 60019620-0202
CURRENT APPLICATION NUMBER: US/10/272,411
CURRENT FILING DATE: 2002-10-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/727,963A
CURRENT FILING DATE: 2000-12-01
PRIOR APPLICATION NUMBER: U.S.S.N. 60/327,557
PRIOR FILING DATE: 1999-12-02
NUMBER OF SEQ ID NOS: 99
                                                                                                                                                                                     NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/329,393 PRIOR FILING DATE: 2001-10-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence: Peptide Ligand NAME/KEY: VARIANT LOCATION: (1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                ENGTH:
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   Conservative
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                     85.7%;
66.7%;
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 0; Mismatches
                   Score 12; DB 14;
Pred. No. 1.4e+06;
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Pred. No. 1.4e+0
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JS-10-181-937-57

OTHER INFORMATION: Description of Artificial Sequence:mp97 and hp97 OTHER INFORMATION: homology consensus peptide

ORGANISM: Artificial Sequence FEATURE:

Matches Query Match Best Local

l Similarity 2; Conserva

85.7%; 66.7%;

Score 12; DB 15; Pred. No. 1.4e+06; 0; Mismatches 1

Length 5; Indels

0

0;

Conservative

1 TXY 3

SEQ ID NO 57

SOFTWARE:

PatentIn Ver. 2.1

PRIOR APPLICATION NUMBER: US 60/181,091
PRIOR FILING DATE: 2000-02-08
PRIOR APPLICATION NUMBER: WO PCT/CA01/00133
PRIOR FILING DATE: 2001-02-08
NUMBER OF SEQ ID NOS: 103

CURRENT APPLICATION NUMBER: US/10/181,937
CURRENT FILING DATE: 2003-03-03

US-09-854-799-40

Sequence 40, Application U Patent No. US20020115064A1 GENERAL INFORMATION:

Application US/09854799

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APPLICANT: Kahan Leong
APPLICANT: Arthur B. Raitano
APPLICANT: Arthur B. Raitano
APPLICANT: Louglas C. Saffran
TITLE OF INVENTION: NOVEL 13-TRANSMEMBRANE PROTEIN EXPRESSED
TITLE OF INVENTION: IN PROSTATE CANCER
FILE REFERENCE: 51158-70011.10
CURRENT APPLICATION NUMBER: US/10/285,045
CURRENT FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: US/9/547,789
PRIOR FILING DATE: 2000-04-12
PRIOR APPLICATION NUMBER: 60/128,858
PRIOR FILING DATE: 1999-04-12
INUMBER OF SEQ ID NOS: 71
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Barnes Jewish Hospital
APPLICANT: Lam, Jonathan
APPLICANT: Rose, F. Patrick
APPLICANT: Teitelbaum, Steven
TITLE OF INVENTION: RANKL MIMICS AND USES THEREOF
FILE REFERENCE: 60019620-0206
CURRENT APPLICATION NUMBER: US/10/272,328A
CURRENT FILING DATE: 2003-01-24
PRIOR APPLICATION NUMBER: 60/329,393
PRIOR FILING DATE: 2001-10-15
NUMBER OF SEQ ID NOS: 51
NUMBER OF SEQ ID NOS: 51
RESULT 11
US-10-284-660-61
; Sequence 61, Application US/10284660
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                                                                                                                                                                                                                                                      ; TYPE: PRT ; ORGANISM: Homo sapiens US-10-285-045-61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; TYPE: PRT; ORGANISM: Homo sapiens US-10-272-328A-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-10-272-328A-26
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Best Local Similarity
Matches 2; Conserv
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn version 3.1
SEQ ID NO 26
LENGTH: 6
                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                              SEQ ID NO 61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Daniel E.H. Afar
APPLICANT: Rene S. Hubert
APPLICANT: Kahan Leong
                                                                                                                                                                       Local Similarity nes 2; Conserv
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Pred. No. 1.4e+06;
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FILE REFERENCE: 51158-20011.01
CURRENT APPLICATION NUMBER: US/10/284,660
CURRENT FILING DATE: 2002-10-30
PRIOR APPLICATION NUMBER: US 09/547,789
PRIOR FILING DATE: 2000-04-12
PRIOR APPLICATION NUMBER: 60/128,858
PRIOR APPLICATION NUMBER: 60/128,858
PRIOR FILING DATE: 1999-04-12
NUMBER OF SEQ ID NOS: 71
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 61
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Best Local Similarity
Watches 2; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            밁
                                                                                                                      RESULT 13
US-10-284-569-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-304-059-42; Sequence 42, Application US/10304059; Publication No. US20030158141A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; ORGANISM: Homo sapiens
US-10+284-660-61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 12
GENERAL INFORMATION:
APPLICANT: Jabbour, Henry Nicolas
APPLICANT: Sales, Kurt Jason
APPLICANT: Katz, Arieh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: GROMEIER, Matthias
APPLICANT: DOBRIKOVA, Elena
                                                                       Sequence 7, Application US/10284569
Publication No. US20030220266A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 42
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/304,059
CURRENT FILING DATE: 2003-03-28
PRIOR APPLICATION NUMBER: 60/332,554
PRIOR FILING DATE: 2001-11-26
NUMBER OF SEQ ID NOS: 47
SOFTMARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: GENETICALLY STABLE EXPRESSION VECTOR FILE REFERENCE: 1579-769
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Kahan Leong
APPLICANT: Arthur B. Raitano
APPLICANT: Douglas C. Saffran
TITLE OF INVENTION: NOVEL 13-TRANSMEMBRANE PROTEIN EXPRESSED
TITLE OF INVENTION: IN PROSTATE CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Daniel E.H. Afar APPLICANT: Rene S. Hubert
                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH:
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2; Conservative
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 Mismatches

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Pred. No. 1.4e+06;
                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                 Score 12; DB 14;
Pred. No. 1.4e+06;
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RESULY 17
US-10-646-874-40
; Sequence 40, Application US/10646874
; Publication No. US20040063093A1
; GENERAL INFORMATION:
APPLICANT: SmithKline Beecham, Corporation
; TITLE OF INVENTION: Recombinant Feline Coronavirus
proteins
                                                                 US-10-646-874-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           밁
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Query Match
Best Local Similarity
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Best Local (
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CURRENT FILING DATE: 2002-10-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 30 SOFTWARE: SeqWin99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Method of treating a disease FILE REFERENCE: ARDW/P27354US
                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 40: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 19406-2799

ZIP: 19406-2799

COMPUTER READABLE FORM:

MEDILM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/646,874

FILING DATE: 22-Aug-2003

PRIOR APPLICATION: 435

PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ENGTH:
                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/392,459
FILING DATE: 22-FEB-1995
APPLICATION NUMBER: US 07/698,927
FILING DATE: 13-MAY-1991
APPLICATION NUMBER: US 07/613,066
FILING DATE: 14-MOV-1990
ATTORNEY/AGENT INFORMATION:
NAME: King, William T.
REGISTRATION NUMBER: 30,954
REFERENCE/DOCKET NUMBER: 30,954
REFERENCE/DOCKET NUMBER: SBC 14532B
TELECOMMUNICATION INFORMATION:
                                                                          TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKliné Beecham Corporation
STREET: 709 Swedeland Road
CITY: King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 TSY 3
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                                                                                                                                                                   LENGTH: 6 amino acids
                                                                                                                                                                                                                                TELEPHONE: (215) 270-5015
TELEFAX: (215) 270-5090
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85.7%;
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Score 12; DB 15; Length 6; Pred. No. 1.4e+06;
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Pred. No. 1.4e+06;
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Search completed: December 13, 2004, 23:02:41 Job time : 69.6 secs
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                                                                                                                                                                    ; TYPE: amino acid
; STRANDEDNESS: unknowr
; TOPOLOGY: unknown
; MOLECULE TYPE: protein
US-08-701-278-14
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US-08-701-278-14
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NAME: Silva, Robin M.
REGISTRATION NUMBER: 38,304
REFERENCE/DOCKET NUMBER: A-63
TELECOMMUNICATION INFORMATION:
TELEPRONE: (415) 781-1989
TELEPRAX: (415) 398-3249
INFORMATION FOR SEQ ID NO: 14:
                                                                                         Query Match
Best Local Similarity 65.,
Thes 2; Conservative
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Publication No. US20030099938A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 7 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert
STREET: Four Emparcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Anderson, David J.
APPLICANT: Saito, Tetsuichiro
TITLE OF INVENTION: A NOVEL HOMEODOMAIN PROTEIN
NUMBER OF SEQUENCES: 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
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Pred. No. 1.4e+06
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2: pir2:*
3: pir3:*
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Copyright (c) 1993 - 2004 Compugen Ltd.
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PQ0150	S22605	S57233	S57230	T14568	T09121	S58569	S05124	H86822	C95215	D64117	D82251	S23365	S6644B	PH0249	B81006
dnaK-type molecula	ribosomal protein	sex combs reduced	(1	hypothetical prote	ftsH protein homol	hypothetical prote	T-cell receptor al	trimethylamine deh	T-cell receptor Vb	hypothetical prote					

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RESULT 3
B39111
Ig heavy chain V region - Pacific hagfish (fragment) C;Species: Eptatretus stouti (Pacific hagfish)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R;Komatsu, S.; Furukawa, K.; Abe, K.; Hirano, H.; Ueda, M. Chem. Pharm. Bull. 40, 2873-2875, 1992
A;Title: Isolation and characterization of equinatoxins from the sea anemone Actinia eq A;Reference number: PC1149; MUID:93099631; PMID:1361161
A;Accession: PC1149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       equinatoxin 1A - sea anemone (Actinia equina) (fragment) (;Species: Actinia equina equi
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(;Species: Arabidopsis thaliana (mouse-ear cress)
(;Date: 30-Jun-1992 #sequence_revision 06-Jan-1995 #text_change 23-Mar-1995
(;Accession: PA0037
R;Kamo, M.; Kawakami, T.; Miyatake, N.; Tsugita, A.
submitted to JIPID, July 1994
A;Description: Separation and characterization of Arabidopsis proteins by tw
A;Reference number: PA0001
A;Accession: PA0037
A;Accession: PA0037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ś
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A; Residues: 1-12 < KAM>
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Best Local Similarity 66.7
Matches 2; Conservative
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Best Local Similarity
Matches 2; Conserv
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66.7%; Pred. No. 8.9e+02;
tive 0; Mismatches 1
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Pred. No. 9.6e+02;
0; Mismatches 1
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R;Levinson, D.A.; Campos-Torres, J.; Leder, P.
J. Exp. Med. 178, 317-329, 1993
A;Title: Molecular characterization of transgene-induced A;Reference number: PH1580; MUID:93301609; PMID:8315387
A;Accession: PH1630
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Date: 18-Oct-1991 #sequence_revision 18-Oct-1991 #text_change (;Accession: B39111  
R;Varner, U; Neame, P.; Litman, G.W. R;Varner, U; Neame, P.; Litman, G.W. 1746-1750, 1991  
proc. Natl. Acad. Sci. U.S.A. 88, 1746-1750, 1991  
A;Title: A serum heterodimer from hagfish (Eptatretus stoutii)  
A;Reference number: A39111; MUID:91156684; PMID:2000382  
A;Accession: B39111  
A;Status: preliminary  
A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                     A43826
dhaK-Type molecular chaperone - Ajellomyces capsulata (fragment)
N;Alternate names: 80K heat shock protein 70 homolog
C;Species: Ajellomyces capsulata, Histoplasma capsulatum
C;Date: 22-Jan-1993 #sequence_revision 22-Jan-1993 #text_change
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                                                                                                                                                                                                                                                                                                                                 R;Gomez, F.J.; Gomez, A.M.; Deepe Jr., Infect. Immun. 60, 2565-2571, 1992 A;Title: An 80-kilodalton antigen from
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C; Keywords: immunoglobulin
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A;Residues: 1-17 <-tEV>
A;Experimental source: bone marrow pre-B lymphocyte
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C;Keywords: heterotetramer; immunoglobulin
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,Molecule type: protein
,,Residues: 1-18 <GOM>
,,Residues: UNIPROT:Q7M4V0
,,Cross-references: UNIPROT:Q7M4V0
,and the control of cell wall and
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;Molecule type: protein
;Residues: 1-14 <VAR>
                                                                                                                                                                                                                                                                                                        Title: An 80-kilodalton antigen from Histoplasma capsulatum that has homology to heat Reference number: A43826; MUID:92307875; PMID:1612726
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         g H chain V-D-J region (clone B-less 156) - mouse (fragment)
;Species: Mus musculus (house mouse)
;Date: 02-Jun-1994 #sequence_revision 02-Jun-1994 #text_change 17-Mar-1999
;Accession: PH1630
                                                                                                                                             , Description: involved in protein folding and assembling/disassembling of , Superfamily: heat shock protein 70\,
                                                                                                                                                                                                                                                                                         Accession: A43826
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Best Local
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T-cell surface glycoprotein RTa - rabbit (fragment)
N,Alternate names: thymocyte specific class I-like antigen, heavy chain
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 09-Jul-2004
C;Pacession: PL0025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Biochem. J. 316, 893-900, 1996
A,Title: Purification and characterization of a tetrameric alpha-macroglobulin proteina
A,Reference number: S70612; MUID:98265058; PMID:8670168
                                                                                                                               T cell receptor alpha chain V region (clone 2DN V alpha 24-4) - C;Species: Homo sapiens (man)
C;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change
C;Accession: PH1783
R;Porcelli, S.; Yockey, C.E.; Brenner, M.B.; Balk, S.P.
J. Exp. Med. 178, 1-16, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: UNIPROT:Q7M3F4
A;Experimental source: thymus
C;Keywords: glycoprotein; surface
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R;Wang, C.R.; Chen, G.H.; Newkirk, M.; Capra, J.D.; Mandy, W.J. Mol. Immunol. 25, 945-952, 1988
A;Title: Biochemical properties of a novel rabbit thymocyte specific class A;Reference number: PL0025; MUID:89096985; PMID:3264885
A;Accession: PL0025
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PL0025
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                                                     A;Reference number: PH1754; MUID:93301585; A;Accession: PH1783
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                                                                                                               J. Exp. Med. 178, 1-16, A; Title: Analysis of T
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A; Residues: 1-18 <BEN>
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A;Residues:
                A; Molecule type:
                                   A; Status: preliminary
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Matches
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1.3e+03;
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RESULT 11
B42382
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$12391

C;Species: Escherichia coli plasmid pFK99 (fragment)
C;Species: Escherichia coli
C;Date: 19-Mar-1997 #sequence_revision 25-Apr-1997 #text_change 09-Jul-2004
C;Accession: $1391
C;Accession: $1391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 homeotic protein Hox-1 - rat (fragment)
(;Species: Rattus norvegicus (Norway rat)
(;Date: 27-Feb-1997 #sequence_revision 27-Feb-1997 #text_change 16-Aug-2004
(;Accession: 165242
R;Sakoyama, Y:; Mizuta, I:; Ogasawara, N:; Yoshikawa, H:
Blochem. Genet. 32, 351-360, 1994
A;Title: Cloning of rat homeobox genes.
A;Reference number: 152340; MUID:95217128; PMID:7702549
A;Accession: 165242
replication initiation protein RepA - Escherichia coli plasmid pMU720 (fragment) (Species: Escherichia coli (C;Decies: Escherichia coli C;Date: 10-Jul-1992 #sequence_revision 10-Jul-1992 #text_change 17-Feb-1994 (C;Accession: B42382 R;Praszkier, J.; Wilson, I.W.; Pittard, A.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Gene: Hox-A; Hox-1
C; Superfamily: homeobox P
C; Keywords: DNA binding;
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A;Molecule type: DNA
A;Residues: 1-20 <RES>
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A,Title: Structure, localization and function of Fanf, a minor component of K99 fibrilla A,Reference number: S12391; MJID:91211613; PMID:1982454
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C;Genetics:
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A; Residues: 1-22 <SIM>
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Pred. No. 1.5e+03;
0; Mismatches 1;
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Pred. No. 1.4e+03;
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J. Bacteriol. 174, 2376-2383, 1992
A;Title: Mutations affecting translational coupling between the rep genes of an IncB mi A;Reference number: A42382; MUID:92202169; PMID:1372603
A;Accession: B42382
                                                                                                                      A;Molecule type: mRNA
A;Residues: 1-24 <MCHS
A;Residues: 1-25 <MCHS
A;Residues: 1-26 <MCHS
A;Experimental source: B cell
A;Note: the authors translated the codon GTA for residue 11 as Thr
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                        C;Accession: PH1712

R;MCHeyzer-Williams, M.G.; McLean, M.J.; Lalor, P.A.; Nossal, G.J.V.
J. Exp. Med. 178, 295-307, 1993

A;Title: Antigen-driven B cell differentiation in vivo.
A;Reference number: PH1675; MUID:93301607; PMID:8315385

A;Accession: PH1712
                                                                                                                                                                                                                                                                                                                                                                                        Ig heavy chain V region (clone ASC-10) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 24-Feb-1994 #sequence_revision 24-Feb-1994 #text_change 17-Mar-1999
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J. Exp. Med. 178, 295-307, 1993
A;Tile: Antigen-driven B cell differentiation in vivo.
A;Reference number: PH1675; MUID:93301607; PMID:8315385
A;Accession: PH1713
A;McJecule type: mENA
A;Residues: 1-24 <MCH>
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C;Species: Mus musculus (house mouse)
C;Date: 24-Feb-1994 #sequence_revision 24-Feb-1994 #text_change 17-Mar-1999
C;Accession: PH1713
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C;Genetics:
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A; Residues: 1-23 < PRA>
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                                                             Score 12; DB 2;
Pred. No. 1.7e+03;
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Pred. No. 1.7e+03;
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Pred. No. 1.6e+03;
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13 TAY 15

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A;Molecule type: protein
A;Residues: 1-25 <OLI>
A;Residues: 1-25 <OLI>
C;Superfamily: omega-conotoxin
C;Keywords: acetylcholine release inhibition; amidated carboxyl end; calcium channel inh
F;1-16,8-20,15-25/Disulfide bonds: #status predicted
F;25/Modified site: amidated carboxyl end (Cys) #status predicted
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C;Species: Electrophorus electricus (electric eel)
C;Date: 31-Mar-1988 #sequence_revision 31-Mar-1988 #text_change 09-Jul-2004
C;Accession: B27262
R;Conti-Tronconi, B.M.; Hunkapiller, M.W.; Lindstrom, J.M.; Raftery, M.A.
Proc. Natl. Acad. Sci. U.S.A. 79, 6489-6493, 1982
A;Title: Subunit structure of the acetylcholine receptor from Electrophorus electricus.
A;Reference number: A93932; MUID:83065199; PMID:6959131
A;Accession: B27262
Search completed: December 13, 2004, 22:33:01 Job time : 17.2 secs
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A;Residues: 1-25 <HIL>
A;Residues: 1-25 <HIL
A;Residues: 1-25 <HIL
A;Residues: 1-25 <HIL
A;Reference number: A34115; MUID:87299637; PMID:2441741
A;Accession: B34115
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C;Species: Conus magus cone)
C;Species: Conus magus (magus cone)
C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 09-Jul-2004
C;Accession: JH0701; B34115
R;Hillyard, D.R.; Monje, V.D.; Mintz, I.M.; Bean, B.P.; Nadasdi, L.; Ramachandran, J.; |
Neuron 9, 69-77, 1992
A;Title: A new conus peptide ligand for mammalian presynaptic Ca2+ channels.
A;Reference number: JH0699; MUID:92337922; PMID:1352996
A;Status: nucleic acid sequence not shown
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A;Residues: 1-24 <CON>
A;Cross-references: UNIPROT:P09689
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Q0s800 saccharomyc
Q6jw56 trochodendr
Q6zy44 silene invo
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                                                    Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids; lamiids; Lamiales; Lamiaceae; Nepetoideae; Ocimeae; Plectranthus. NCBI TaxID=204187;
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NON_TER
SEQUENCE
    Paton A.,
                SEQUENCE FROM N.A.
PubMed=15019625;
                                                                                                                          Plectranthus glabratus.
                                                                                                                                             Name=rps16
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Springate D.A., Sudde S., Otieno
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ALIGNMENTS

Q9BYF9 PRELIMINARY; Q9BYF9; 01-JUN-2001 (TrEMBLrel. 17, C 01-JUN-2001 (TrEMBLrel. 17, L 01-JUN-2003 (TrEMBLrel. 24, L Cyrokeratin 19 (Fragment). Kagaya M., Kaneko S., Ohno H., Inamura K., Kobayashi K.; "Cloning and characterization of the 5'-flanking region of human cytokeratin 19 gene in human cholangiocarcinoma cell line."; J. Hepatol. 35:504-511(2001). SEQUENCE FROM N.A. MEDLINE=21539745; PubMed=11682035; Kagaya M., Kaneko S., Ohno H., Ina Homo sapiens (Human). Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. EMBL; AB045973; BAB40770.1; -. GO; GO:0005882; C:intermediate filament; Conservative 9 1122 MW; 85.7%; 66.7%; Score 12; DB 2 Pred. No. 1.8e-0; Mismatches 0 Created) Last sequence update) Last annotation update) 9E9FC41B45AB45A1 CRC64; 1.8e+06; DB 2; A 1; Indels Length 9; 0; Gaps 0

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                                                                                                            Q94IS6 PRELIMINARY; PRT; 10 AA. Q94IS6 (94IS6)
01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last amotation update)
Nonspecific lipid transfer protein (Fragment).
Pinus taeda (Lobbolly pine).
Pinus taeda (Lobbolly pine).
Eukaryota; Viridiplantae; Streptophyta; Embryophy
Spermatophyta; Coniferopsida; Coniferales; Pinace
  Submitted (MAR-2001) to the EMBL/GenBank/DDBJ databases
                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Paton A., Springate D.A., Sudde S., Otieno D., Willis F., Simonds M.S.J., Powell M.P., Savola "Phylogeny and evolution of basils and allies based on three plastid DNA regions."; Mol. Phylogenet. Evol. 31:277-299(2004).

EMBL; AJ505387; CAD45507.1; -.
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EMBL; AJ505387; CAD45507.1; -.

EO; GO:0003735; F:structural constituent of ribosome; IEA.
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14-MAR-2004 (TrEMBLrel. 27, Created)
14-MAR-2004 (TrEMBLrel. 27, Last sequence update)
14-MAR-2004 (TrEMBLrel. 27, Last annotation update)
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lamiids; Lamiales; Lamiaceae; Nepetoideae; Ocimeae; Plectranthus.
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PubMed=15019625;
Paton A., Springate D.A., Sudde S., Otieno D.,
Paton A., Springate D.A., Sudde S., Otieno D.,
Willis F., Simonds M.S.J., Powell M.P., Savola'
"Phylogeny and evolution of basils and allies
"Phylogeny and evolution of basils and allies
based on three plastid DNA regions.";
Mol. Phylogenet. Evol. 31:277-299(2004).
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NON TER
SEQUENCE
Paton A., Springate D.A., Sudde S., Otieno D., Grayer R., Harley M.M., Willis F., Simonds M.S.J., Powell M.P., Savolainen V.; "Phylogeny and evolution of basils and allies (Ocimeae, Labiatae) based on three plastid DNA regions.";
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                                                                                                       SEQUENCE FROM N.A.
                                                                                                                                                                               Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids; Lamiids; Lamiales; Lamiaceae; Nepetoideae; Ocimeae; Ocimum.
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14-MAR-2004 (TrEMBLrel. 27, Last seg
14-MAR-2004 (TrEMBLrel. 27, Last ann
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
                                                                                                                                                         NCBI_TaxID=204098;
                                                                                                                                                                                                                                                                                                                                           Ribosomal protein
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L; AJ505420; CAD45540.1; -.
GO:0003735; F:structural constituent
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Pred. No. 3.8e+0
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Pred. No. 3
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3.2e+03;
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Savolainen V.;
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Q7BVK6;
05-UUL-2004 (TrEMBLrel. 2
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Streptomycin resistance p
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila; mayaguana subcluster.
NCBI TaxID=214822;
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EMBL, AJ505420; CAD45540.1; -
Chloroplast; Ribosomal protein.
NON TER 1 1 1 1 NON TER 12 12
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Submitted (SEP-2002) to the EMBL/GenBank/DDBJ databases
EMBL; AY154986; AA013634 1; -.
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MEDLINE=98027397; PubMed=9361446; Palmer E.L., Teviotdale B.L., Jones A.L.; "A relative of the broad-host-range plasmid RSF1010 Erwinia amylovora."; Appl. Environ. Microbiol. 63:4604-4607(1997). EMBL; AF017390; AAC45878.1; -.
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                                                                                                                                                   NCBI_TaxID=552;
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NGRAD P.M. II, Durando C.M., Heed W.B.,
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nes 2; Conserv
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protein B
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Pred. No. 3.8e+03;
0; Mismatches 1;
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Pred. No. 4.1e+03;
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01-JUN-2002
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                                                                                              Name=tnpA;
Pseudomonas fluorescens.
Bacteria; Proteobacteria;
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Submitted (MAR-2004) to the EMBL/GenBank/DDBJ
EMBL; AJ634219; CAG24647.1; -.
GO; GO:0009507; C:chloroplast; IEA.
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Spermatophyta; Magnoliophyta; eudicotyledons;
Caryophyllales; Caryophyllaceae; Silene.
NCBI_TaxID=269082;
SEQUENCE FROM N.A.
STRAIN=TC29-5;
MEDLINE=22315381; PubMed=12427948;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Disentangling the reticulate history of polyploids in Silene (Caryophyllaceae).";
Thesis (2004), Department of Evolutionary Biology, Systematic
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05-UUL-2004 (TrEMBLrel. 27, Last sequence update)
05-UUL-2004 (TrEMBLrel. 27, Last annotation update)
Cytochrome b6/f complex subunit VI (Fragment).
                                                                                Pseudomonadaceae; Pseudomonas.
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                                                                 NCBI_TaxID=294;
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TISSUE=Leaf;
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Uppsala,
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Pred. No. 4.1e
0; Mismatches
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Pred. No.
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CAG24647
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P80625;
01-OCT-1996
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CAG24647;
31-MAR-2004
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NCBI_TaxID=269082;
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"Tn5041-like transposons: molecular diversity, evolutionary
relationships and distribution of distinct variants in environmental
bacteria.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Popp M., Erixon P., Eggens F., Oxelman B.; "Oxigin and evolution of a circumpolar polyploid species complex in Silene (Caryophyllaceae)."; Submitted (MAR-2004) to the EMBL/GenBank/DDBJ databases.
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31-MAR-2004 (TrEMBLrel. 27, Last sequence update)
31-MAR-2004 (TrEMBLrel. 27, Last annotation update)
31-MAR-2004 (TrEMBLrel. 27, Last annotation update)
Cytochrome b6/f complex subunit VI (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                              Chloroplast.
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EMBL; AJ634219; CAG24647.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Thesis (2004), Department of Evolutionary 
Uppsala university, Uppsala, Sweden.
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(Rel. 34, Created)
(Rel. 34, Last seq
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Pred. No. 4.4e
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Q6ZYT6;
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05-JUL-2004
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Zea mays (Maize).

Bukaryota, Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;

PACCAD clade; Panicoideae; Andropogoneae; Zea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lee M.-H., Schedl T.;
Submitted (OCT-2003) to the
EMBL; AY455927; AAR26306.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Touzet P., Riccardi F., Morin C., Damerval C., Huet J.-C., pernollet J.-C., Zivy M., de Vienne D.; "The maize two dimensional gel protein database: towards an integrated genome analysis program."; Theor. Appl. Genet. 93:997-1005(1996). "Theor. Appl. Genet. 93:997-1005(1996). "Theor. Appl. is: 5.6, its MW is: 18.4 kDa. Maize-2DPAGE; P80625; COLEOFTILE.
  Silene
                 Cytochrome b6/f complex subunit VI (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Caenorhabditis elegans.
Eukaryota; Metazoa; Nematoda; Chromadorea;
Rhabditidae; Peloderinae; Caenorhabditis.
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05-JUL-2004
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05-JUL-2004
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Unknown protein from 2D-PAGE of etiolated coleoptile (Spot 406)
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Pred. No. 4
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Pred. No. 4.7e+03;
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Lee M.-H., Schedl T.;
Lee M.-H., Schedl T.;
Lee m.-H., Schedl T.;
"Translation repression by GLD-1 protects its mRNA targets from non-
sense mediated mRNA decay.";
Submitted (CCT-2003) to the EMBL/GenBank/DDBJ databases.

EMBL; AY455927; AAR26306.1;
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SEQUENCE 15 AA; 1901 MW; 86D1593EFA18C498 CRC64;
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Submitted (MAR-2004) to the EMBL/GenBank/DDBJ databases.
EMBL; AJ634220; CaG24449-1; -
GO; GO:0009507; C:chloroplast; IEA.
Chloroplast.
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SEQUENCE 15 AA; 1645 MW; C716428D1B688740 CRC64;
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Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Rhabditoidea;
Rhabditidae; Peloderinae; Caenorhabditis.
NCBI_TaxID=6239;
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Chloroplast.
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Caryophyllales; Caryophyllaceae; Silene.
NCBI_TaxID=269082;
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Thesis (2004), Department of Evolutionary Biology, Systematic Botany,
Uppsala university, Uppsala, Sweden.
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